



(12) **United States Patent**
O'Shea et al.

(10) **Patent No.:** **US 7,902,348 B2**
(45) **Date of Patent:** **Mar. 8, 2011**

(54) **JANUS FAMILY KINASES AND IDENTIFICATION OF IMMUNE MODULATORS**

(75) Inventors: **John J. O'Shea**, Silver Spring, MD (US); **Warren J. Leonard**, Bethesda, MD (US); **James A. Johnston**, Middletown, MD (US); **Sarah M. Russell**, Kensington, MD (US); **Daniel W. McVicar**, Charles Town, WV (US); **Masaru Kawamura**, Rockville, MD (US)

(73) Assignee: **The United States of America as represented by the Department of Health and Human Services**, Washington, DC (US)

(*) Notice: Subject to any disclaimer, the term of this patent is extended or adjusted under 35 U.S.C. 154(b) by 0 days.

(21) Appl. No.: **12/354,668**

(22) Filed: **Jan. 15, 2009**

(65) **Prior Publication Data**
US 2009/0137787 A1 May 28, 2009

Related U.S. Application Data

(60) Division of application No. 11/195,197, filed on Aug. 1, 2005, now Pat. No. 7,488,808, which is a continuation of application No. 08/373,934, filed on Jan. 13, 1995, now Pat. No. 7,070,972.

(51) **Int. Cl.**
C12N 15/52 (2006.01)
C12N 15/12 (2006.01)

(52) **U.S. Cl.** **536/23.2**

(58) **Field of Classification Search** None
See application file for complete search history.

(56) **References Cited**

U.S. PATENT DOCUMENTS

5,124,251	A	6/1992	Lanier et al.
5,262,522	A	11/1993	Gearing
5,283,354	A	2/1994	Lemischka
5,705,625	A	1/1998	Civin et al.
5,916,792	A	6/1999	Civin et al.
6,136,595	A	10/2000	Ihle et al.

FOREIGN PATENT DOCUMENTS

WO 94/00469 1/1994

OTHER PUBLICATIONS

Asao, et al. 1994. "Interleukin 2-induced activation of JAK3: possible involvement in signal transduction for c-myc induction and cell proliferation." *FEBS Lett.* 351:201-206.
Beadling, et al. 1994. "Activation of JAL kinases and STAT proteins by interleukin-2 and interferon α , but not the T cell antigen receptor, in human T lymphocytes." *EMBO J.* 13:5605-5615.
Bennett, et al. 1994. "Identification and characterization of a novel tyrosine kinase from megakaryocytes." *J. Biol. Chem.* 269:1068-1074.

Cance, et al. 1993. "Novel protein kinases expressed in human breast cancer." *Int. J. Cancer* 54:571-577.

Dyrberg, et al. 1986. "Peptides as antigens." *J. Exp. Med.* 164:1344-1349.

Fasciglione, et al. 1996. "Hapten-carrier interactions and their role in the production of monoclonal antibodies against hydrophobic haptens." *Hybroma* 15(1):1-9.

Gnarra, et al. 1990. "Human interleukin 2 receptor β -chain gene: chromosomal localization and identification of 5' regulatory sequences." *PNAS USA* 87:3440-3444.

Hunter, T. 1993. "Cytokine Connections." *Nature* 366:114-116.

Johnston, et al. 1994. "Phosphorylation and activation of the Jak-3 Janus kinase in response to interleukin-2." *Nature* 370:151-153.

Kawamura, et al. 1994. "Molecular cloning of L-JAK, a Janus family protein-tyrosine kinase expressed in natural killer cells and activated leukocytes." *PNAS USA* 91:6374-6378.

Kirken, et al. 1993. "Characterization of an interleukin-2 (IL-2)-induced tyrosine phosphorylated 116-kDa protein associated with IL-2 receptor β -subunit." *J. Biol. Chem.* 268:22765-22770.

Lai, et al. 1994. "Characterization and expression of human JAK3 splice variants." *Blood* 84 Suppl. 1, 294A.

McVicar, et al. 1994. "Molecular Cloning of *lsk*, a carboxyl-terminal *src* kinase (*csk*) related gene, expressed in leukocytes." *Oncogene* 9:2037-2044.

Miyazaki, et al. 1994. "Funktional activation of Jak1 and Jak3 by selective association with IL-2 receptor subunits." *Science* 266:1045-1047.

Nakamura, et al. 1994. "Heterodimerization of the IL-2 receptor β and γ chain cytoplasmic domains is required for signalling." *Nature* 369:330-333.

Noguchi, et al. 1993. "Characterization of the human interleukin-2 receptor γ chain gene." *J. Biol. Chem.* 268:13601-13608.

Rane, et al. 1994. "JAK3: a novel JAK kinase associated with terminal differentiation of hematopoietic cells." *Oncogene* 9:2415-2423.

Russell, et al. 1994. "Interaction of IL-2 receptor β chain and γ_c chains with JAK1 and JAK3, respectively: defective γ_c -JAK3 association in X-linked severe combined immunodeficiency." *Aids Res. Human Retrovir.* 10:S73.

Russell, et al. 1993. "Interleukin-2 receptor γ chain: a functional component of the interleukin-4 receptor." *Science* 262:1880-1883.

Russell, et al. 1994. "Interaction of IL-2R β and γ_c chains with Jak1 and Jak3: implications for XSCID and XCID." *Science* 266:1042-1045.

Safford, et al. 1994. "JAK3, a member of the JAK family of non-receptor tyrosine kinases, is expressed in the stem/progenitor cell fraction of human bone marrow." *Blood* 84 Suppl. 1, 122A.

Sanchez, et al. 1994. "Multiple tyrosine protein kinases in rat hippocampal neurons: isolation of Ptk-3, a receptor expressed in proliferative zones of the developing brain." *PNAS USA* 91:1819-1823.

(Continued)

Primary Examiner — Ron Schwadron

(74) *Attorney, Agent, or Firm* — Swanson & Bratschun, L.L.C.

(57) **ABSTRACT**

An isolated polynucleotide encodes JAK-3 protein. JAK-3 protein is a protein tyrosine kinase having a molecular weight of approximately 125 kDa which has tandem non-identical catalytic domains, lacks SH2 or SH3 domains, and is expressed in NK cells and stimulated or transformed T cells, but not in resting T cells. The protein itself and antibodies to this protein are also presented. Further, methods of identifying therapeutic agents for modulating the immune system make use of the foregoing.

2 Claims, No Drawings

OTHER PUBLICATIONS

Sharon, et al. 1990. "A 100-kilodalton protein is associated with the murine interleukin 2 receptor: biochemical evidence that p100 is distinct from the α and β chains." *PNAS USA* 87:4869-4873.

Siliciano, et al. 1992. "*itk*, a T-cell-specific tyrosine kinase gene inducible by interleukin 2." *PNAS USA* 89:11194-11198.

Stahl, et al. 1993. "The alphas, betas, and kinases of cytokine receptor complexes." *Cell* 74:587-590.

Takahashi, et al. 1994. "Molecular cloning of rat JAK3, a novel member of the JAK family of protein tyrosine kinases." *FEBS Lett* 342:124-128.

Takebe, et al., 1988. "Src α a promoter: an efficient and versatile mammalian cDNA expression system composed of the simian virus 40 early promoter and the R-U5 segment of human T-cell leukemia virus type 1 long terminal repeat." *Mol. Cell. Biol.* 8:466-472.

Vitte-Mony, et al. 1994. "Signal transduction of interleukin 2 in human natural killer cells: involvement of the p56^{lek} tyrosine kinase." *Mol. Immunol.* 31:623-632.

Withuhn, et al. 1994. "Involvement of the Jak-3 Janus kinase in signalling by interleukins 2 and 4 in lymphoid and myeloid cells." *Nature* 370:153-157.

Zeng, et al. 1994. "JAK3 Janus kinase is involved in interleukin 7 signal pathway." *FEBS Lett.* 353:289-293.

1

JANUS FAMILY KINASES AND IDENTIFICATION OF IMMUNE MODULATORS

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is a divisional of U.S. patent application Ser. No. 11/195,197, filed Aug. 1, 2005, now issued as U.S. Pat. No. 7,488,808, which is a continuation of U.S. application Ser. No. 08/373,934, filed Jan. 13, 1995, now issued as U.S. Pat. No. 7,070,972, both of which are hereby expressly incorporated by reference in their entireties.

SEQUENCE LISTING

The present application is being filed along with a Sequence Listing in electronic format. The Sequence Listing is provided as a file entitled NIH100.1C1DV1.TXT, created Dec. 23, 2008, which is 27 Kb in size. The information in the electronic format of the Sequence Listing is incorporated herein by reference in its entirety.

FIELD OF THE INVENTION

The present invention relates generally to the field of transmembrane signal transduction. More specifically, the invention relates to the components of multisubunit receptors for cytokines, and assays for detecting interactions between these components.

BACKGROUND OF THE INVENTION

Protein tyrosine kinases (PTK) are critical enzymes for receptor-mediated signal transduction in lymphocytes. Indeed, protein tyrosine phosphorylation is an early and requisite event in signaling for both multichain immune recognition receptors such as the T-cell antigen receptor (TCR) and cytokine receptors. Unlike growth factor receptors, neither the TCR nor the cytokine receptors have intrinsic PTK activity. Rather, they are coupled to nonreceptor tyrosine kinases. For example, the src family PTK, Lck and Fyn have been implicated in TCR-mediated signaling. The non-src family PTK Zap-70 has been shown to associate with the TCR upon activation. The src family PTK, Lck, Fyn and Lyn have also been implicated in Interleukin-2 (IL-2) receptor mediated signaling.

The Janus family kinases (JAKs) represent a recently described family of PTK. These kinases, called JAK-1, JAK-2 and Tyk-2, are structurally quite distinct in that they possess tandem nonidentical catalytic domains. These PTK are ubiquitously expressed and have been shown to participate in signaling by a number of cytokine and hormone receptors. These PTK are believed to exert their effects through tyrosyl phosphorylated transcription factors.

IL-2 is a T-cell lymphokine that both functions as a potent autocrine growth factor and activates other cells, including B-cells and natural killer (NK) cells. In addition, the interaction of IL-2 with high affinity IL-2 receptors regulates the magnitude and duration of the normal T-cell immune response.

High affinity IL-2 receptors comprise three receptor components, denoted the IL-2 receptor α , β and the common γ_c chain. Interestingly, the γ_c chain is a receptor component that is shared by the receptors for several of the interleukins. In the mechanism of IL-2 signal transduction, IL-2 binding induces

2

heterodimerization of the cytoplasmic domains of β and γ_c . This heterodimerization is required for IL-2 signaling.

While the IL-2 receptor is a multi-subunit complex that lacks intrinsic enzymatic activity, it is clear that protein tyrosine phosphorylation is an early biochemical step that follows ligand binding. Occupancy of the IL-2 receptor additionally stimulates alkalization of T-cells via Na^+/H^+ exchange, activation of Ras, Raf and ERK1, induction of the fos, jun and myc proto-oncogenes, as well as induction of effector functions such as cytotoxicity and cellular proliferation.

SUMMARY OF THE INVENTION

One aspect of the invention provides an isolated polynucleotide encoding the JAK-3 protein. The JAK-3 protein is a protein tyrosine kinase that has a molecular weight of approximately 125 kDa. In addition, the JAK-3 protein is characterized by tandem non-identical catalytic domains. The JAK-3 protein lacks SH2 or SH3 domains, and is expressed in NK cells and stimulated or transformed T-cells, but not substantially expressed in resting T-cells. In a preferred embodiment, the isolated polynucleotide encoding the JAK-3 protein has the nucleotide sequence of SEQ ID NO:8.

A second aspect of the invention relates to an isolated JAK-3 protein. The JAK-3 protein is a protein tyrosine kinase having a molecular weight of approximately 125 kDa, has tandem non-identical catalytic domains, lacks SH2 or SH3 domains, and is expressed in NK cells and stimulated or transformed T-cells, but not substantially expressed in resting T-cells. In a preferred embodiment, the isolated JAK-3 protein has the polypeptide sequence of SEQ ID NO:9.

A third aspect of the invention relates to an antibody that is specific to JAK-3 protein. As described above, the JAK-3 protein is a protein tyrosine kinase having a molecular weight of approximately 125 kDa. The JAK-3 protein recognized by the claimed antibody has tandem non-identical catalytic domains, lacks SH2 or SH3 domains, and is expressed in NK cells and stimulated or transformed T-cells, but not substantially expressed in resting T-cells. In a preferred embodiment, the antibody that specifically recognizes the JAK-3 protein is polyclonal.

A fourth aspect of the invention provides a method of identifying an agent having immunomodulating activity. This method includes first determining the ability of the β or γ_c chain of the IL-2 receptor to physically associate with a Janus family kinase selected from the group consisting of JAK-1 and JAK-3 in the absence of a candidate immunomodulatory agent. A second step includes determining the ability of a chain of the IL-2 receptor to physically associate with a Janus family kinase in the presence of the candidate immunomodulating agent. If the candidate immunomodulating agent causes either a lesser or greater ability of the IL-2 receptor chain and the Janus family kinase to associate with each other, then such an ability will indicate that the candidate agent has immunomodulating activity. In a preferred embodiment, the two determining steps are performed in the presence of a cytokine compound. In a more preferred embodiment, the cytokine compound is a cytokine compound that binds to a receptor that includes the γ_c chain. In yet a more preferred embodiment, the cytokine compound is selected from the group consisting of IL-2, IL-4, IL-7, IL-9 and IL-15. In another preferred embodiment, each of the determining steps includes immunoprecipitating a chain of the IL-2 receptor with an antibody that is specific to that chain, and then determining if the Janus family kinase coprecipitates with that

receptor chain. According to this embodiment, the candidate agent can be a peptide fragment of the γ_c chain of the IL-2 receptor.

A fifth aspect of the invention relates to a method of identifying a therapeutic agent for modulating the immune system. This method includes first isolating a Janus family kinase selected from the group consisting of JAK-1 and JAK-3. A subsequent step involves determining the ability of the kinase to phosphorylate a substrate in the absence of a candidate agent, and then determining the ability of the kinase to phosphorylate the substrate in the presence of the candidate agent. A decrease in the ability of the kinase to phosphorylate the substrate in the presence of the candidate agent indicates that said candidate agent is an immunosuppressive agent. Alternatively, a greater ability of the kinase to phosphorylate the substrate in the presence of the candidate agent indicates that said candidate agent is an immuno-enhancing agent. In a preferred embodiment, the isolating step includes immunoprecipitating the Janus family kinase with an antibody specific to the Janus family kinase. In a more preferred embodiment, each of the determining steps includes first reacting the Janus family kinase with γ -labeled ATP to produce reaction products, and then separating the reaction products by electrophoresis and identifying the labeled products. In an even more preferred embodiment, each of the determining steps includes first reacting the Janus family kinase with ATP to produce reaction products, then separating the reaction products by electrophoresis, exposing the separated reaction products to a labeled antibody specific for phosphotyrosine, and finally identifying the reaction products that exhibit binding to the antibody.

DETAILED DESCRIPTION OF THE INVENTION

We have discovered that a novel protein tyrosine kinase (PTK), called JAK-3, has structural features characteristic of the Janus family of PTK. Unlike the three other members of this family, JAK-3 is only expressed in a narrow spectrum of cell types. Whereas JAK-1, JAK-2 and Tyk2 are broadly expressed, JAK-3 is the first Janus family member that exhibits limited tissue expression, or that is induced following cellular activation. In particular, JAK-3 was found to be predominantly expressed in activated human T lymphocytes and NK cells. We also have observed JAK-3 expression in IFN- γ activated human peripheral blood monocytes and B-cells. Based on this pattern of gene expression, we believe that the function of JAK-3 is related to important signaling pathways in these cell types. However, we acknowledge that JAK-3 could also be inducible in nonhematopoietic tissues by appropriate stimuli.

Herein we disclose that IL-2 can rapidly induce activation and tyrosine phosphorylation of JAK-3 in responsive cells. Additionally, we show that JAK-1, although less prominently tyrosine-phosphorylated, also is involved in IL-2 receptor signaling. The data presented below indicate that both JAK-3 and JAK-1 play important roles in lymphoid activation via the IL-2 receptor. Specifically, JAK-3 and JAK-1 associate with other components of the IL-2 receptor to form an activated receptor complex. Thus, agents that inhibit the interaction of either JAK-1 or JAK-3 with other components of the receptor are considered as candidates for immunomodulatory drugs. We contemplate that such immunomodulatory drugs can be either immunosuppressing or immunoenhancing.

More specifically, compounds that inhibit the interaction of JAK-1 with IL-2R β , or the interaction of JAK-3 with the γ_c subunit of the IL-2 receptor complex are expected to inhibit IL-2 dependent signal transduction. These inhibitory com-

pounds can act through mechanisms involving drug interaction with the kinase, or with the receptor substrate. In the first instance, the compound may interact with the kinase such that the kinase fails to phosphorylate the receptor chain. Alternatively, the inhibitor may act on the kinase such that it fails to autophosphorylate. According to another contemplated mechanism, the inhibitory compound may interact with one of the chains of a multisubunit receptor, so that interaction with the Janus family kinase is inhibited. Finally, we contemplate compounds which disrupt the binding between one receptor chain and the Janus family kinase that ordinarily binds to that receptor subunit.

The development of assays for identifying compounds that inhibit IL-2 dependent signalling represents an important step toward the discovery of novel immunomodulators. Compounds having inhibitor activity in such assays will be identified as potential therapeutics for the treatment of autoimmune diseases and transplant rejection. Autoimmune diseases anticipated as objects of therapy using drugs identified according to the assays disclosed herein include, but are not limited to, rheumatoid arthritis, psoriatic arthritis, lupus and vasculitis. Certain of the drugs identified according to the present invention are likely to be useful not only in the treatment of disorders resulting from inappropriate activation of the IL-2 receptor, but also from inappropriate activation of other cytokine receptors that share common subunits with the IL-2 receptor.

Our approach to the development of an assay for compounds that inhibit IL-2 dependent signalling stems from our understanding of the detailed structure of the IL-2 receptor complex. In particular, we have discovered that two different Janus family tyrosine kinases interact with different subunits of the IL-2 receptor. Binding of the JAKs to the receptor subunits represents a critical step in the cytokine signalling pathway. It follows that any agent that disrupts or prevents this binding will also inhibit IL-2 dependent signalling. Thus, we have designed an assay to identify compounds that inhibit interactions between components of the IL-2 receptor complex as a method for identifying candidate agents for modulating the immune system.

Direct physical interactions of IL-2R β with JAK-1 and γ_c with JAK-3 were demonstrated during the development of the present invention. IL-2 increased JAK-3 association with the IL-2 receptor, with an apparent increase of γ_c -JAK-3 association and de novo induction of IL-2R β -JAK-3 interaction. Truncation of γ_c resulted in its failure to associate with JAK-3. Moreover, a patient having a mild form of X-linked combined immunodeficiency (XCID) that was characterized by diminished IL-2 induced proliferation had a γ_c point mutation (Leu 271 changed to Gln) that decreased γ_c association with JAK-3. Thus, γ_c mutations, in at least some XSCID and XCID patients, can inhibit the ability of γ_c to associate with JAK-3. The severity of this immunodeficiency owes to the fact that γ_c is a component of multiple cytokine receptors.

Indeed, the sharing of the γ_c subunit of the IL-2 receptor by other cytokine receptors, including receptors for IL-4, IL-7, IL-9 and IL-15, has led us to believe that any compound which inhibits the cytokine induced binding of JAK-3 to the γ_c chain would also inhibit signalling through the receptors for any of these cytokines. Thus, we believe that an assay for compounds that inhibit cytokine induced binding of JAK-3 to the γ_c chain is useful in the discovery of drugs that could be used to treat autoimmune diseases that are attributable to the activities of IL-2, IL-4, IL-7, IL-9 and IL-15.

Although other materials and methods similar or equivalent to those described herein can be used in the practice or testing of the present invention, the preferred methods and

5

materials are now described. General references for methods that can be used to perform the various PCR and cloning procedures and nucleic acid and protein blotting procedures described herein can be found in *Molecular Cloning: A Laboratory Manual* (Sambrook et al. eds. Cold Spring Harbor Lab Publ. 1989) and *Current Protocols in Molecular Biology* (Ausubel et al. eds., Greene Publishing Associates and Wiley-Interscience 1987). The disclosures contained in these publications are hereby incorporated by reference. A description of the experiments and results that led to the creation of the present invention follows.

We employed the PCR approach described by Harpur et al., in *Oncogene* 7:1347 (1992) and by Siliciano et al., in *Proc. Natl. Acad. Sci. USA* 89:11194 (1992) to amplify novel PTK encoding sequences expressed in NK cells. The disclosures of these Harpur et al. and Siliciano et al. references are incorporated by reference. Briefly, we first prepared cDNA from NK cell mRNA using reverse transcriptase. We then performed PCR using degenerate oligonucleotide primers corresponding to conserved motifs in the catalytic domains of PTK. The forward primers were designed to correspond to residues in subdomain VI and to exclude src-family PTK. The reverse primer corresponded to the reverse complement of the DVWSFG (SEQ ID NO:1) motif (subdomain IX) conserved in a large number of PTK.

Example 1 describes the methods used to isolate an amplified polynucleotide that corresponded to the JAK-3 tyrosine kinase.

EXAMPLE 1

PCR Amplification of a Novel NK Cell Derived Protein Tyrosine Kinase

First strand cDNA was prepared from total RNA that had been isolated from NK cell RNA according to standard methods. The cDNA prepared in this fashion served as the template in a subsequent PCR reaction.

The forward primer used in this PCR reaction was 5'-CCA GCG GCC GCG T(G/A/T/C)CA (C/T)CG (G/A/T/C)GA (C/T)C T(G/A/T/C)GC-3' (SEQ ID NO:2) and the reverse primer was 5'-CCA GCG GCC GCC C(G/A)A A(G/A/T/C/)(G/C) (A/T)CC A(G/A/T/C)A C(G/A)T C-3' (SEQ ID NO:3). The resulting products were digested with NotI, subcloned and sequenced. The PCR fragment corresponding to one novel kinase was isolated, labeled and used to screen several cDNA libraries. Among the libraries screened were oligo (dT) primed cDNA libraries derived from PHA stimulated peripheral blood T-cells and the HUT-78 T-cell line (Clonetechn, Palo Alto, Calif.) in the λ gt11 cloning vector, a λ ZAP YT library and a λ ZAP library from PHA activated T-cells. Purified phage DNA was digested and subcloned into pBLUESCRIPT (Stratagene, La Jolla, Calif.) for nucleic acid sequencing. Sequence data were manipulated and analyzed using the programs of the Genetics Computer Group of the University of Wisconsin and the BLAST program of NCBI.

For Northern analysis, total RNA from various human tissues was either purchased from Clonetechn (Palo Alto, Calif.) or prepared from NK cells and T-cells. Northern blotting was carried out according to standard procedures. Blots were probed with radiolabeled nucleic acid probes corresponding to the PCR amplified fragment described above.

Results indicated that one of the candidate clones isolated by this method was preferentially expressed in NK and activated T-cells as an approximately 4.3 kb mRNA.

Thus, the DNA fragment amplified using kinase-specific primers exhibited a tissue-specific pattern of mRNA expres-

6

sion. We therefore proceeded to isolate a cDNA clone that contained the entire reading frame for the protein that was partly encoded by the amplified fragment. We have termed this protein "JAK-3."

Example 2 describes the method used to determine the DNA sequence of the polynucleotide that encoded the putative tissue-restricted kinase obtained in Example 1.

EXAMPLE 2

Isolation and Analysis of a Polynucleotide Encoding JAK-3

The PCR generated fragment from Example 1 was used as a probe to screen phage cDNA libraries prepared from the NK-like cell line, YT, PHA-activated T-cells and HUT-78 cells. Approximately 5×10^5 plaques from each library were screened to obtain multiple overlapping clones that generated a polynucleotide sequence corresponding to a single long open reading frame. SEQ ID NO:9 provides the predicted amino acid sequence encoded by the NK cell derived JAK-3 cDNA. We note that a published report by Kawamura et al. (*Proc. Natl. Acad. Sci. USA* 91:6374 (1994)) refers to the JAK-3 protein disclosed herein as "L-JAK." The polynucleotide sequence encoding the JAK-3 protein has been filed in GenBank under accession number U09607 and is provided herein as SEQ ID NO:8.

The open-reading frame of this polynucleotide sequence encompasses 3372 nucleotides and is predicted to encode a polypeptide having 1124 residues. The molecular weight of the predicted polypeptide is 125,014 Da. This value is roughly equivalent to, but slightly smaller than, other JAKs. As described below, this predicted molecular weight is consistent with the M_r of the polypeptide identified by Western blotting using the antiserum prepared against JAK-3.

The predicted protein encoded by the JAK-3 polynucleotide exhibited structural features characteristic of a PTK, as would readily be appreciated by one having ordinary skill in the art. Like other PTK, a catalytic domain is present in the C-terminal portion of the molecule (subdomains I-XI). This domain begins with a typical ATP-binding motif at residues 829-834 (subdomain I) in which the canonical GXGXXG (SEQ ID NO:4) motif is evident and is followed by a critical lysine residue in subdomain II (residue 855). Just C-terminal to subdomain VII is a pair of tyrosine residues following an acidic residue that could represent the autophosphorylation site. In subdomain VIII, phenylalanine and tyrosine residues surround the invariant tryptophan residue. We note that the locations of the subdomains cited herein are identified by Kawamura et al. in *Proc. Natl. Acad. Sci. USA* 91:6374 (1994), the disclosure of which is hereby incorporated by reference. This atypical motif contrasts with the motifs seen in src and abl related proteins, and growth factor receptors. Notably though, this motif (FWYAPE) (SEQ ID NO:5) is present in the Janus family kinases. The Janus family of PTK comprises PTK that have tandem nonidentical catalytic domains and a large extracatalytic segment. These structural features of the Janus family of PTK have been considered by Wilks et al., in *Mol Cell. Biol.* 11:2057 (1991), Harpur et al., in *Oncogene* 7:1347 (1992), and Firmbach-Kraft et al., in *Oncogene* 5:1329 (1990). The entire catalytic domain, termed the JAK homology (JH) 1 domain, comprises 273 amino acids (residues 822 to 1095) and is followed by a unique C-terminus.

In addition to a kinase catalytic domain, the predicted JAK-3 protein has a region N-terminal to the PTK catalytic domain that also has elements typical of a protein kinase

catalytic domain. This tandem kinase-like (JH-2) domain is a characteristic feature of the JAKs, and has been described by Harpur et al., in *Oncogene* 7:1347 (1992), Firmbach-Kraft et al., in *Oncogene* 5:1329 (1990), Velazquez et al., in *Cell* 70:313 (1992), and Argetsinger et al., *Cell* 74:237 (1993). Like other Janus family kinases, this domain in the NK derived JAK-3 protein lacks some standard features of a PTK catalytic domain. In particular, the JAK-3 protein appears to lack an autophosphorylation site in subdomain VII.

The known JAK family PTK have large extracatalytic segments (JH 3-7 domains) located N-terminal to the kinase (JH1) and kinase-like (JH2) domains. While motifs corresponding to SH2 or SH3 domains are lacking, there is a motif that Harpur et al., (*Oncogene* 7:1347 (1992)) suggested to have SH2-like character. This region is conserved in three of the four family members, including the JAK-3 protein. Immediately N-terminal to this motif is a highly conserved motif that is a possible tyrosine phosphorylation site (VDGYFRL) (SEQ ID NO:6). Other areas of striking homology between the novel JAK-3 protein and other JAKs are also evident in the remaining domains (JH 5-7).

The findings presented above indicated the NK-derived isolate had nearly all of the characteristics of a JAK family PTK. Protein homology analysis indicated the overall identity of the NK PTK to the most closely related Janus family member, JAK-2, was approximately 68%. The fact that a hydrophilicity plot showed no evidence for a hydrophobic domain suggested that the JAK-3 polynucleotide sequence encoded a non-receptor type PTK.

Other members of the Janus family of PTK (JAK-1, JAK-2, and Tyk2) have been shown to be present in a variety of tissues. These findings have been described by Wilks et al., in *Mol. Cell. Biol.* 11:2057 (1991), Harpur et al., in *Oncogene* 7:1347 (1992), and Firmbach-Kraft et al., in *Oncogene* 5:1329 (1990). Given this precedent, we proceeded to determine if JAK-3 was also broadly expressed across different tissues and cell types. As described below, we unexpectedly discovered that JAK-3 expression is tissue-restricted.

Example 3 describes the methods used to determine the expression profile for the JAK-3 mRNA.

EXAMPLE 3

Expression of the JAK-3 mRNA

Total RNA (20 µg) from various human tissues (purchased from Clontech), NK cells and T-cells was electrophoresed in formaldehyde/agarose gels, transferred to membranes and probed with a radiolabeled cDNA corresponding to the JH1 and JH2 domains of JAK-3. T-cells were activated with PHA for 24 hours prior to RNA isolation. NK cells were activated with IL-2 (1,000 units/ml) for 24 hours prior to RNA isolation. Uniform RNA loading in the different lanes of the Northern blot was confirmed by ethidium bromide staining and by hybridization with a probe that detected ribosomal RNA.

Unexpectedly, and in contrast to other Janus family members, JAK-3 exhibited a restricted pattern of mRNA expression across different tissues. In the absence of stimulation, the mRNA was only detected in NK cells. Notably, activation of NK cells did not alter the level of JAK-3 mRNA expression. The JAK-3 mRNA was also detected at high levels in activated NK cells and activated T-cells. No JAK-3 mRNA was detected in liver, testis, kidney, small intestine, brain or lung tissues. Hybridization of the same Northern Filter with a JAK-1 cDNA probe gave evidence for mRNA expression in all tissues except for small intestine. Interestingly, while the

JAK-3 mRNA was expressed at very low levels in resting T-cells, the mRNA was induced upon T-cell activation.

Various cultured cell lines were also tested for expression of the JAK-3 mRNA. As expected, our results indicated the JAK-3 mRNA was constitutively expressed in the NK-like YT cell line. As predicted by studies with peripheral blood T-cells, the JAK-3 mRNA was not constitutively expressed in the Jurkat T-cell line but was induced upon activation. Interestingly, we found that JAK-3 mRNA was constitutively expressed in the HUT-78 transformed T-cell line. No expression of this gene was detected in a variety of other cell lines, including the erythroleukemia cell line, K562.

We next identified a unique portion of the C-terminal region of the JAK-3 protein and generated a corresponding synthetic peptide for use as an immunogen. The conclusion regarding protein sequence uniqueness was based on the results from a computer-assisted comparison between the JAK-3 amino acid sequence and all sequences available through the Program Manual for the Wisconsin Package, Version 8, September 1994, Genetics Computer Group, (575 Science Dr. Madison, Wis. 53711). This search protocol included searches of the GenBank, GenPept, SwissProt, Brookhaven and EMBL databases. The synthetic peptide used in our procedures corresponded to amino acids 1104-1124 of the JAK-3 protein.

Example 4 describes the immunological methods used to analyze the JAK-3 protein.

Example 4

Identification of the JAK-3 Protein by Metabolic Labeling and Immunoprecipitation

A peptide corresponding to the predicted C-terminus of the JAK-3 protein (amino acids 1104-1124) was synthesized (Multiple Peptide Systems, San Diego, Calif.), coupled to keyhole limpet hemocyanin (KLH) with MBS (Pierce, Rockford, Ill.) and used to immunize rabbits. HUT-78 cells (10^7 cells per point) were labeled with ^{35}S methionine (0.5 mCi/ml) for 2 hours, washed with phosphate buffered saline and lysed in buffer containing 1% TRITON™ X-100 detergent (lysis buffer). Postnuclear supernatants were immunoprecipitated with 10 µl of antiserum prebound to protein A sepharose washed in buffer containing 0.1% TRITON™ X-100 detergent (wash buffer), eluted and electrophoresed in 8% polyacrylamide gels that were subsequently fixed, rinsed in FLUORO-HANCE™ fluorographic enhancer (Research Products Inc., Mount Prospect, Ill.) and dried for autoradiography.

For immunoblot analysis, cells were solubilized in lysis buffer and postnuclear supernatants (approximately 100 µg of protein) were electrophoresed, transferred to nitrocellulose and immunoblotted. Filters were blocked, incubated with antiserum (1:1000), washed and incubated with peroxidase conjugated goat antirabbit IgG, all according to standard methods well known in the art. Antibody binding was detected by enhanced chemiluminescence (ECL) (Amersham Corp.).

In good agreement with the molecular weight predicted by the deduced JAK-3 primary structure, analysis of metabolically labeled HUT-78 cells showed specific immunoprecipitation of a polypeptide with a M_r of approximately 125 kDa. Immunoblot analysis of these cells also showed reactivity of the antibody with a protein of approximately the same mobility in HUT-78. In contrast Jurkat T-cells expressed minimal levels of this protein. In additional experiments, the expression of the JAK-3 encoded polypeptide was found to parallel

the expression seen by analysis of mRNA. Expression of the protein was detected in NK cells, activated T-cells and in some transformed leukocyte cell lines. Immunoblotting with preimmune serum or antiserum competed with cognate peptide versus irrelevant peptide, thus confirming the specificity of this reactivity.

The results obtained in Example 4 confirmed that antibodies raised against a JAK-3 specific synthetic peptide could be used to immunoprecipitate the JAK-3 protein from cellular lysates. However, these results provided no insight into the function of the JAK-3 protein. Thus, to ascertain whether the JAK-3 protein had enzymatic activity, in vitro kinase assays were performed.

Example 5 describes the in vitro assay used to demonstrate that immunoprecipitated JAK-3 protein had kinase activity.

EXAMPLE 5

JAK-3 In Vitro Kinase Activity

Kinase assays were performed as described by Muller et al., in *Nature* 366:129 (1993), and by Watling et al., in *Nature* 366:166 (1993). According to these procedures, cells were solubilized in lysis buffer supplemented with 1 mM Na_3VO_4 and 1 mM EDTA. An antipeptide antiserum was then used to carry out an immunoprecipitation reaction. Washed immunoprecipitates were incubated in 50 μl of buffer containing 20 mM Tris, 5 mM MgCl_2 , 5 mM MnCl , 1 μM ATP, and 200 $\mu\text{Ci/ml}$ ^{32}P -ATP (Amersham Corp.). The reaction was carried out for 15 minutes at 25° C. and was terminated by the addition of ice cold wash buffer. After washing the beads again, the reaction products were eluted, electrophoresed and autoradiographed.

A phosphorylated polypeptide having the expected 125 kDa molecular weight was evident in immunoprecipitates from NK cells but not in immunoprecipitates from resting T-cells or untreated control cells. The phosphorylated residues were resistant to KOH, consistent with tyrosyl phosphorylation. We concluded that this tyrosyl phosphorylation likely represented autophosphorylation of JAK-3.

Thus, our results indicated the novel JAK-3 protein had both structural and functional characteristics of a protein tyrosine kinase related to the Janus family of PTK. In an effort to discern the role of JAK-3 in cellular physiology, we made a more extensive investigation into the range of cell types that expressed the JAK-3 protein.

Example 6 describes the method used to determine the range of cell types that express JAK-3 protein.

EXAMPLE 6

Expression of the JAK-3 Protein in Activated T-Cells and NK Cells

Whole cell lysates from human peripheral blood T-cells (unstimulated or stimulated for 48 hours with PHA), the transformed T-cell lines Hut78 and YT, peripheral blood NK cells, the NK 3.3 cell line, human peripheral blood monocytes, the myelomonocytic cell lines U937 and THP-1, and the tumor cell line OVCAR-3, HT-29 and IM-9 were run on SDS-PAGE and were probed with antisera to JAK-3. Human peripheral blood T lymphocytes, NK cells and monocytes (>97% pure) were obtained by leukopheresis and column purification according to standard procedures. T lymphocytes were either untreated or treated with PHA (10 $\mu\text{g/ml}$) and incubated for 0-48 hours. Cells were grown in RPMI 1640 supplemented with calf serum. Cells were lysed in buffer

containing TRITON™ X-100 detergent and clarified lysates (50 μg) were run on SDS PAGE, transferred to IMMOBILON™ membrane and probed with anti-JAK-3 antisera and HRP-conjugated antirabbit immunoglobulin. Immunoblots were developed with enhanced chemiluminescence (Amersham) using standard procedures. The anti-JAK-3 antiserum used in these procedures was raised against a synthetic peptide corresponding to the C-terminal region of the JAK-3 protein (amino acids 1104-1124), and has been described by Kawamura et al., in *Proc. Natl. Acad. Sci. USA* 91:6374 (1994).

The results from these procedures indicated that JAK-3 was expressed only in a limited spectrum of tissue types. It was strongly expressed in NK cells, the NK cell line NK 3.3, YT cells and in the transformed T-cell line Hut78 as judged by the presence of a 125 kDa band on the Western blot. The JAK-3 protein was not detected in other cell types that were examined. In contrast to the pattern of JAK-3 protein expression, other JAK family kinases are known to be expressed in both lymphoid and nonlymphoid cells. While JAK-3 was expressed at low levels in resting peripheral blood T-cells, these levels of expression were greatly increased following activation by either PHA or anti-CD3.

The induction of the IL-2 gene and IL-2 receptor α chain are among the critical events that occur during T-cell activation. These factors have been considered by Taniguchi in *Ann. Rev. Immunol.* 4:69 (1988), and by Leonard et al., in *Proc. Natl. Acad. Sci. USA* 80:6957 (1983). As the induction of these genes paralleled the induction of JAK-3, we investigated the possibility that JAK-3 was somehow coupled to the IL-2 receptor. Indeed, in *Cell* 74:587 (1993), Stahl et al. disclosed that occupancy of certain other cytokine receptors induced tyrosine phosphorylation of other JAKs.

Example 7 describes the methods used to demonstrate that the JAK-3 protein is inducibly phosphorylated by IL-2 stimulation of T and NK cell lines, and peripheral blood NK cells.

EXAMPLE 7

Tyrosine Phosphorylation of JAK-3 in Response to IL-2

Cells were either unstimulated or stimulated with IL-2 (1000 U/ml) for 5 minutes, then lysed, immunoprecipitated with anti-JAK-3 antisera and immunoblotted and probed with a monoclonal antiphosphotyrosine antibody (4G10 UBI). In this procedure, cells were washed twice with RPMI that had been acidified at pH 6.5 before incubation for 3 hours with 0.5% human serum. The cells were again washed twice before stimulation with IL-2. The cells were lysed in buffer containing TRITON™ X-100 detergent prior to the immunoprecipitation procedure. Detection of the antiphosphotyrosine antibody probe bound to the 125 kDa JAK-3 protein band was by autoradiography.

Using this procedure, we found that IL-2 induced tyrosine phosphorylation of JAK-3 in YT (T-cell line) and NK 3.3 (NK cell line) cells as well as in peripheral blood NK cells. We observed little basal tyrosine phosphorylation of JAK-3 in any of the cell samples that were tested. However, after 5 minutes of IL-2 stimulation, intense phosphorylation of a protein band having a molecular weight of 125 kDa was evident on the autoradiograph. This band represented the JAK-3 protein. A constitutively expressed phosphoprotein band that migrated at a discrete position in the gel between the 69 and 96 kDa molecular weight markers was observed in YT and NK3.3 cell lines, and in peripheral blood NK cells. A second constitutive phosphoprotein band was observed in the

11

NK3.3 lanes on the autoradiograph. In aggregate, these results indicated that JAK-3 is a protein that is inducibly phosphorylated by stimulation of T-cells and NK cells with IL-2.

IL-2 induced tyrosine phosphorylation in the Hut78 T-cell line was also examined in a time course experiment. Lysates from Hut78 cells either untreated or treated with IL-2 for 5, 10 or 30 minutes were immunoprecipitated with anti-JAK-3 and blotted with antiphosphotyrosine antibody and with anti-JAK-3. Our results indicated that Hut78 cells showed induction of JAK-3 tyrosine phosphorylation after treatment with IL-2 that was maximal at 10 minutes. Prior absorbance of the JAK-3 antiserum with cognate peptide eliminated the tyrosine phosphorylated protein at 125 kDa and thereby confirmed the specificity of the tyrosine phosphorylated band as JAK-3. JAK-3 was also found to be phosphorylated in response to IL-2 in human peripheral blood T-cells that had been pre-activated with PHA for 24 hours. Thus in human T and NK cells, T-cell and NK cell lines, IL-2 induced the rapid tyrosyl phosphorylation of JAK-3 protein.

Both Farrar, et al. (*J. Biol. Chem.* 264:12,562 (1989)) and Kirken, et al., (*J. Biol. Chem.* 268:22, 765 (1993)) have disclosed that IL-2 stimulation induced tyrosyl phosphorylation of a variety of substrates in human peripheral blood T-cells and T-cell lines. In the following Example, we demonstrated that the most prominent phosphotyrosyl protein evident following IL-2 stimulation of the YT cell line was a polypeptide of 125 kDa that comigrated with JAK-3. To confirm the identity of this 125 kDa substrate as JAK-3, an experiment was conducted in which lysates from IL-2 treated cells were first depleted of JAK-3 prior to immunoprecipitation with antiphosphotyrosine antibodies. In addition, we examined whether the other JAK family members could be activated by IL-2.

Example 8 describes the methods used to demonstrate that JAK-1 and JAK-3 are tyrosine phosphorylated in T-cells upon IL-2 stimulation.

EXAMPLE 8

IL-2 Induced Tyrosyl Phosphorylation of JAK Family Kinases

Serum-starved YT cells were treated with IL-2 (1000 U/ml) for 0 or 15 minutes. Cells were lysed in buffer containing TRITON™ X-100 detergent and phosphatase inhibitors and then immunoprecipitated with an antiphosphotyrosine antibody bound to protein G sepharose. Lysates were pre-cleared with either rabbit polyclonal antisera, anti-JAK-3 or anti-JAK-1 before immunoprecipitation and immunoblotting with anti-phosphotyrosine. As a positive control to identify the position of JAK-3 on the Western blot, a sample of the lysate from IL-2 induced cells was immunoprecipitated with anti-JAK-3 and blotted with antiphosphotyrosine. All immunoprecipitates were subjected to SDS-PAGE and transferred to IMMOBILON™ membrane.

Depletion of YT cell lysates with anti-JAK-3 antiserum specifically removed the 125 kDa tyrosine phosphorylated protein, thus indicating that JAK-3 was one of the most prominent phosphoproteins detected in response to IL-2 in these cells. Conversely, depletion of lysates with anti-JAK-1 did not remove this phosphoprotein. This result clearly established that JAK-3 is a prominent tyrosyl phosphoprotein in IL-2 induced T-cells.

In addition, we examined whether other JAK family members could be activated by IL-2. An experiment similar to that described above was performed in which antiphosphoty-

12

rosine immunoblots were purposely overexposed to reveal other substrates. In particular, YT cells were treated with IL-2 (1000 U/ml) for 0, 5 or 15 minutes, lysed, immunoprecipitated with anti-phosphotyrosine and subsequently immunoblotted with anti-phosphotyrosine, and anti-JAK-3 or anti-JAK-1. We observed a tyrosyl phosphoprotein induced in response to IL-2 that was larger in size than JAK-3. Immunoprecipitation with anti-JAK-1 antiserum and immunoblotting with antiphosphotyrosine confirmed that this protein was JAK-1. Although both Tyk2 and JAK-2 could be detected in YT cells, no tyrosyl phosphorylation of either protein was observed in response to IL-2 stimulation.

Thus, the results obtained in Example 8 indicated that IL-2 stimulation of T-cells led to tyrosine phosphorylation of JAK-3 and, to a somewhat lesser extent, of JAK-1.

Example 9 describes the methods used to demonstrate that IL-2 stimulates JAK-3 in vitro kinase activity.

EXAMPLE 9

IL-2 Stimulates JAK-3 in vitro Kinase Activity

Cells were treated with IL-2 (1000 U/ml) for 0, 10 or 20 minutes. In vitro kinase activity was measured in anti-JAK-3 immunoprecipitates. For measurement of in vitro kinase activity, immunoprecipitates were washed in 50 mM NaCl, 5 mM MgCl₂, 5 mM MnCl₂, 0.1 mM Na₃VO₄, 10 mM HEPES (pH 7.4) and incubated in the same buffer containing 0.25 μ Ci/ml [³²P- γ] ATP for 15 minutes at room temperature. After three washes, proteins were eluted in sample buffer and analyzed by SDS-PAGE. Gels were either dried or transferred to IMMOBILON™ membrane before exposure. Autophosphorylation signals were detected by autoradiography of the dried gels. JAK-3 in vitro kinase activity in response to IL-2 stimulation (15 minutes) was measured in immunoprecipitates from peripheral blood NK cells and Hut78 cells. JAK-3 immunoprecipitated from YT cells activated in response to IL-2 displayed elevated kinase activity that peaked at approximately 10 minutes. Similar results were obtained using human peripheral blood NK cells and Hut78 cells. This kinase activity was not precipitated by preimmune serum or in the presence of competing JAK-3 peptide. These results confirmed that the observed kinase activity was specific for the JAK-3 protein. The vast majority of the phosphorylation observed was attributable to phosphorylation of tyrosine residues as demonstrated by resistance to KOH treatment and by phosphoamino acid analysis.

We next examined JAK-3 phosphorylation in YT cells following stimulation by a number of these cytokines. Other members of the JAK family have been shown to be activated by several lymphokines including the interferons (IFNs), erythropoietin (EPO), growth hormone (GH) and IL-3 (Witthuhn et al., *Cell* 74:227 (1993); Argetsinger et al., *Cell* 74:237 (1993); Silvennoinen et al., *Proc. Natl. Acad. Sci. USA* 90:8429 (1993)). JAK-3 was tyrosine phosphorylated in response to IL-2, but not in response to GH, IFN- α or IFN- γ in the YT cells. We did detect tyrosine phosphorylation of JAK-2 in response to IFN- γ in these cells, thus confirming the activity of this cytokine. Stimulation of YT cells with IL-3, GM-CSF or EPO also failed to induce JAK-3 phosphorylation. IL-4 has recently been shown to utilize the common γ chain of the IL-2 receptor (γ_c), suggesting that common signaling pathways also may be utilized by these cytokines (see Russell et al., *Science* 262:1880 (1993), and Kondo et al., *Science* 262:1874 (1993)). This knowledge prompted us to examine whether JAK-3 was also activated in response to IL-4.

13

Example 10 describes the methods used to test whether other cytokines, in addition to IL-2, induced JAK-3 tyrosine phosphorylation.

EXAMPLE 10

Phosphorylation of JAK-3 in Response to Cytokines
Other than IL-2

YT cells were serum-starved for 3 hours and acid washed twice before stimulation and immunoprecipitation. NK 3.3 cells were grown for 2 days in 10% lymphocult (Biotest) and 15% fetal bovine serum before being washed and incubated in 2% serum. Commercial polyclonal anti-JAK-1 and JAK-2 antisera were obtained from UBI and Tyk2 polyclonal anti-serum from Santa Cruz Biotechnology.

Lysates of YT cells that were unstimulated or stimulated with IL-2 (1000 U/ml), GH (50 ng/ml), IFN- α (1000 U/ml) or IFN- γ (500 U/ml) for 15 minutes were immunoprecipitated with anti-JAK-3 and blotted with antiphosphotyrosine. Results from this Western blotting experiment proved that JAK-3 was tyrosine phosphorylated in response to IL-2, but not in response to GH, IFN- α or IFN- γ . In addition, JAK-3 was not phosphorylated in response to IL-3, granulocyte macrophage colony stimulating factor (GM-CSF) or erythropoietin, all of which have been shown to activate JAK-2.

In related procedures, NK 3.3 cells that were incubated overnight in 2% serum were stimulated with IL-2 (1000 U/ml), IL-4 (100 U/ml) or IFN- γ (500 U/ml) for 5 or 15 minutes. Cellular lysates were immunoprecipitated with anti-JAK-3 and blotted with antiphosphotyrosine. Results from these procedures indicated that JAK-3 was tyrosine phosphorylated in response to both IL-4 and IL-2 in NK 3.3 cells. These results confirmed that the JAK-3 kinase is involved in signaling pathways other than IL-2.

Given these findings, we proceeded to investigate the detailed mechanism by which the JAK proteins were involved in IL-2 mediated signal transduction. As described in the following Examples, we discovered that two of the IL-2 receptor subunits interacted with the JAKs. In particular, we discovered that the JAK-1 and JAK-3 proteins physically interacted with the IL-2R β and γ_c IL-2 receptor subunits. Further, the γ_c -JAK-3 binding was found to be induced by IL-2 stimulation. Our procedures began with a demonstration that both the JAK-1 and JAK-3 proteins are phosphorylated by IL-2 stimulation.

Example 11 describes methods that can be used to detect phosphorylation of the JAK-1 and JAK-3 proteins.

EXAMPLE 11

IL-2 Stimulation Induces Phosphorylation of JAK-1
and JAK-3

Peripheral blood lymphocytes (PBL) were induced for 15 minutes with IL-2 (1000 U/ml), IL-4 (100 U/ml), IL-7 (100 ng/ml), or IL-9 (100 ng/ml) after phytohemagglutinin (PHA) stimulation. The PBL were activated for 72 hours with PHA, then washed twice at pH 6.5, incubated for 3 hours in medium containing 0.5% human serum and resuspended in medium containing 10% fetal calf serum for one hour. Cells were then lysed and immunoprecipitated with polyclonal antibodies to JAK-1 (UBI) or JAK-3. The anti-JAK-3 antibody used in these procedures has been described by Johnston et al., in *Nature* 370:151 (1994). Phosphotyrosine-containing proteins were detected by immunoblotting with 4G10 (UBI). NK3.3 cells were stimulated with IFN- α (1000 U/ml) or IL-2 (1000

14

U/ml), immunoprecipitated with polyclonal antibodies to Tyk2 (UBI) or JAK-2 (UBI) and immunoblotted with 4G10 mAb to phosphotyrosine (UBI). In one procedure, the lysate was precleared with polyclonal antibody to JAK-3.

Although immunoprecipitation with antibodies to JAK-2 yielded a tyrosine phosphorylated band in response to IL-2, this band migrated faster than the JAK-2 band induced by interferon- γ and was human JAK-3, immunoprecipitated through cross-reactivity with the JAK-2 antiserum. This was demonstrated by elimination of this band by preclearing the lysate with a JAK-3 specific antiserum. In addition to IL-2, we tested three other known γ_c users, IL-4, IL-7 and IL-9, for their abilities to induce the tyrosine phosphorylation of JAK-1 and JAK-3. Each of these cytokines induced tyrosine phosphorylation of both JAK-1 and JAK-3 in Western blotting procedures and activated both JAK-1 and JAK-3 as evaluated by in vitro kinase assays. Thus, cytokines that signal using mechanisms that involve the common γ_c subunit induce phosphorylation of JAK-1 and JAK-3.

Given the essential roles of both IL-2R β and γ_c in IL-2 signal transduction, we investigated the ability of each of these chains to physically associate with JAK-1 and/or JAK-3, the two Janus kinases that were activated and tyrosine phosphorylated in response to IL-2. YT cellular lysates were immunoprecipitated with Mik β 1 (anti-IL-2R β) monoclonal antibody (mAb) or R878 (anti- γ_c) antiserum, followed by Western blotting with JAK-3 and reblotting with JAK-1.

Example 12 describes the methods used to demonstrate that JAK-1 binds the IL-2R β subunit, and that JAK-3 binds the γ_c subunit of the IL-2 receptor.

EXAMPLE 12

Physical Association of JAK-1 and JAK-3 with
Components of the IL-2 Receptor

YT cells were either stimulated or not stimulated with IL-2 and then lysed with 10 mM Tris (pH 7.5) containing 2 mM EDTA, 0.15 M NaCl, 0.875% Brij 96, 0.125% NONIDET P40TM detergent, 0.4 mM sodium vanadate, 1 mM 4-(2-aminoethyl)-benzenesulfonyl fluoride hydrochloride (ICN), 2.5 mM leupeptin, 2.5 mM aprotinin. Immunoprecipitations were performed using Mik β 1 anti-IL-2R β mAb, R878 antiserum to γ_c , or RPC5 (control mAb). The gel was Western blotted sequentially with antisera to JAK-3 and then JAK-1 using ECL. The object of this coprecipitation procedure was to test whether precipitation of one subunit of the IL-2 receptor would also precipitate one of the JAK proteins.

Our results indicated that JAK-1 constitutively associated with IL-2R β , and that the quantitative extent of association did not increase following IL-2 stimulation. JAK-1 did not constitutively associate with γ_c , but after IL-2 stimulation and the induction of association of IL-2R β with γ_c , some JAK-1 coprecipitated with γ_c . Although JAK-3 weakly associated with γ_c in the absence of IL-2, this association increased following IL-2 stimulation. Moreover, after IL-2 stimulation, JAK-3 was readily coprecipitated with anti-IL-2R β mAb Mik β 1 or TU11. This was expected for TU11, which binds to an IL-2R β epitope distinct from the IL-2 binding site and coprecipitates γ_c in the presence but not absence of IL-2, but was unexpected for Mik β 1, which competes for IL-2 binding and cannot coprecipitate γ_c in the presence or absence of IL-2.

These data indicated that IL-2R β primarily associated with JAK-1 and that γ_c associated with JAK-3. However, interactions between IL-2R β and JAK-3 also occurred. Though the basis for the IL-2 induced association of IL-2R β with JAK-3 is unknown, IL-2 induced heterodimerization of IL-2R β and

15

γ_c may juxtapose JAK-3 to IL-2R β , thus facilitating their interaction. Significantly, the association of JAK-3 with γ_c was induced by IL-2.

To further evaluate the association of IL-2R β and γ_c with JAK-1, we transiently transfected COS-7 cells with cDNAs encoding JAK-1 and either IL-2R β or γ_c .

Example 13 describes a cotransfection procedure that confirmed the physical interaction between JAK-1 and IL-2R β . The results of this procedure did not provide evidence for an interaction between JAK-1 and the γ_c receptor subunit.

EXAMPLE 13

Recombinant JAK-1 Associates with Recombinant IL-2R β in Transfected Cells

Cotransfection of COS-7 cells with JAK-1 and either IL-2R β or γ_c expression constructs was followed by immunoprecipitation and Western blotting with Erd A antiserum to IL-2R β , R878 antiserum to γ_c or antiserum to JAK-1. The IL-2R β expression construct used in this procedure consisted of the IL-2R β 1 cDNA insert described by Gnarr et al., in *Proc. Natl. Acad. Sci. USA* 87:3440 (1990), under transcriptional control of the SR α promoter that has been described by Takebe et al., in *Mol. Cell. Biol.* 8:466 (1988). The γ_c expression construct used in this procedure included a γ_c cDNA insert derived from a clone that was isolated from a cDNA library prepared from YT cell mRNA. The 5' end of the γ_c cDNA insert corresponded to position -43 as presented in FIG. 2 by Noguchi et al., in *J. Biol. Chem.* 268:13601 (1993), and extended to the 3' end of the full length γ_c sequence that has been presented by Takeshita et al., in *Science* 257:379 (1992). The JAK-1 expression construct used in this procedure had the JAK-1 cDNA under transcriptional control of the CMV promoter. When JAK-1 and IL-2R β expression constructs were cotransfected, antibodies to either protein coprecipitated the other as evaluated by Western blotting. As expected, the RPC5 control mAb failed to immunoprecipitate material that could be identified by either the anti-JAK-1 or the anti-IL-2R β antiserum on the Western blot. This negative result confirmed the specificity of the blotting procedure. The Gnarr et al., Takebe et al., Noguchi et al., and Takeshita et al. references referred to in this example are all incorporated herein by reference.

Our results also indicated that γ_c and JAK-1 did not associate with each other. This latter finding was evidenced by the inability of anti-JAK-1 to immunoprecipitate material that could be stained on Western blots probed with anti- γ_c antibodies, and by the inability of anti- γ_c to immunoprecipitate material that could be stained on Western blots probed with anti-JAK-1. On the other hand, anti-JAK-1 immunoprecipitated material that was stained by anti-JAK-1, and anti- γ_c antibodies immunoprecipitated material that was stained by anti- γ_c antibodies. This latter observation confirmed the integrity of the reagents used in these procedures. As expected, the RPC5 control mAb failed to immunoprecipitate material that was stained by either anti-JAK1 or anti-IL-2R β antiserum on the Western blot. Although the positive control procedures confirmed our ability to detect all relevant protein species, we did not obtain any evidence for an association between JAK-1 and the γ_c receptor chain. However, these findings did confirm a physical association between JAK-1 and IL-2R β .

We conducted an experiment to determine the relative importance of the cytoplasmic domain of the IL-2R β receptor chain, and to test the importance of the extracellular domain with respect to facilitating the interaction between JAK-1 and

16

the IL-2R β chain. COS-7 cells were cotransfected with JAK-1 and either IL-2R α , chimeric $\alpha/\alpha/\beta$ or chimeric $\alpha/\gamma/\gamma$ expression constructs according to standard protocols. The structures of the chimeric constructs encoding these novel receptor chains has been described by Nakamura et al., in *Nature* 369:330 (1994), the disclosure of which is hereby incorporated by reference. The RPC5 control mAb and anti-Tac mAb to IL-2R α were separately used for immunoprecipitations that were followed by Western blotting with antiserum to JAK-1. Neither anti-Tac nor RPC5 antibodies immunoprecipitated material from lysates of JAK-1 and IL-2R α cotransfectants that was also stained by anti-JAK-1. Thus, there was no evidence that JAK-1 interacted with any portion of the IL-2R α chain. Neither anti-Tac nor RPC5 antibodies immunoprecipitated material from lysates of JAK-1 and chimeric $\alpha/\gamma/\gamma$ cotransfectants that was also stained by anti-JAK-1. Thus, there was no evidence that JAK-1 interacted with the cytoplasmic portion of the γ_c receptor chain. On the other hand, when lysates from cells cotransfected with JAK-1 and chimeric $\alpha/\alpha/\beta$ were immunoprecipitated with anti-Tac and Western blotted, material stained by anti-JAK-1 was detected.

These results established that recombinant components of the IL-2 receptor associate with each other in defined ways. Further, our results confirmed the interaction of the JAK-1 and IL-2R β proteins, and indicated that the cytoplasmic portion of the IL-2R β chain was required for this interaction.

We next used transfected COS-7 cells to evaluate the association of JAK-3 with γ_c mutants in order to identify regions of contact between the JAK-3 and γ_c proteins. These coprecipitation experiments employed two different anti- γ_c antibodies, a monoclonal antibody that recognized the extracellular domain of γ_c , and R878, a polyclonal antiserum that recognized the intracellular domains of γ_c . As described below, we discovered that recombinant γ_c and JAK-3 proteins efficiently interacted with each other. In contrast, two truncated forms of the receptor chain in which either 80 or 48 amino acids had been deleted from the C-terminus (Russell et al. *Science* 262:1880 (1993)) exhibited greatly diminished association with JAK-3. The 48 amino acid truncation is smaller than the truncation found in the XSCID patient with the smallest known naturally occurring γ_c truncation (missing 62 amino acids). Other XSCID patients have been identified with even larger truncations. Thus, the findings presented below confirmed that the inability of JAK-3 to associate with γ_c would be a characteristic of many XSCID patients.

Example 14 describes experiments which demonstrated recombinant γ_c chains that incorporate mutations similar to those found in XSCID patients cannot efficiently associate with JAK-1.

EXAMPLE 14

Mutations in Recombinant γ_c Chains Disrupt JAK-3 Binding

COS-7 cells were transfected with JAK-3 and either wild type γ_c or γ_c - Δ CT, γ_c - Δ SH2, γ_2 -L271Q or with the vector control (pME18S). The γ_c expression construct used in these procedures is described under Example 13. The γ_c - Δ CT and γ_c - Δ SH2 constructs were prepared as described by Russell et al., in *Science* 262:1880 (1993). The γ_c -L271Q expression construct was prepared by using the pAlter-1 Mutagenesis Vector system (Promega), and is essentially identical to the wild type γ_c construct, except for mutation of codon 271 from CTG to CAG. The transfectants were lysed and immunoprecipitated with the extracellular domain-recognizing antibody or R878 anti- γ_c antibodies and Western blotted with JAK-3

antiserum. Expression of transfected wild type and mutant γ_c constructs were similar as determined by flow cytometry. Note that whereas the extracellular domain-recognizing antibody mAb bound to an extracellular epitope and thereby all the mutant forms of γ_c tested, R878 cannot bind γ_c - Δ CT or γ_c - Δ SH2 truncation mutants which lack the critical epitope.

The extracellular domain-recognizing antibody immunoprecipitates from cells cotransfected with JAK-3 and wild type γ_c contained material that was Western blotted with anti-JAK-3. No other cotransfectant gave similar results. Repetition of the procedure using the R878 anti- γ_c immunoprecipitates gave essentially similar results, except that a small amount of anti-JAK-3 staining material was observed in the lane corresponding to the lysate from cells that had been cotransfected with JAK-3 and γ_c -L271Q expression constructs. We noted that the amount of JAK-3 associating with γ_c -L271Q was substantially less than the amount of JAK-3 associating with the wild type γ_c . These results indicated that mutation of the amino acid at position 271 of the γ_c chain inhibited JAK-3 binding. This implicated the region of the γ_c protein that included amino acid position 271 as being critical for interaction with the JAK-3 protein.

Neither the vector control nor any of the γ_c mutants that were tested in our experiments gave any indication for JAK-3 binding. Further, we discovered that the portion of the γ_c chain that included amino acid position 271 represented a critical domain of the protein that was required for JAK-3 binding.

We have recently characterized the genetic defect in a pedigree with an X-linked combined immunodeficiency as a single nucleotide change within the γ_c cytoplasmic domain, resulting in replacing Leu 271 with Gln. As described above, the γ_c -L271Q mutation significantly diminished but did not abrogate association with JAK-3. The finding that the γ_c -L271Q mutant still weakly associated with JAK-3 was consistent with the disease phenotype in this pedigree being less severe than typical XSCIDs. Nevertheless, affected males manifest diminished and retarded development of CD4+ and CD8+T-cells and decreased T-cell responses to mitogens and IL-2.

Interestingly, Leu 271 is not contained within the region deleted in the γ_c - Δ SH2 mutant. We interpret this as indicating that JAK-3 may contact residues both proximal and distal to the deletion point in the γ_c - Δ SH2 construct. An experiment described in Example 15, that was conducted using a 19 amino acid long peptide spanning Leu 271, supports this hypothesis since the peptide only partially inhibited γ_c -JAK-3 coprecipitation, even when present at high molar excess. When IL-2R β and JAK-3 were cotransfected into COS-7 cells, Mik β 1 anti-IL-2R β mAb weakly but reproducibly coprecipitated JAK-3.

As described above, we have disclosed that JAK-3 is indeed the kinase that associates with γ_c and that JAK-3 is activated and tyrosine phosphorylated by all previously known γ_c users tested, as well as IL-9, which we now add to the list of known γ_c users. Moreover, we have partially defined the regions and residues of γ_c required for JAK-3 association and correlated defective γ_c -JAK-3 association with XSCID.

The activation of JAK-1 by IL-2, IL-4, IL-7, and IL-9 was an unexpected result and suggests that IL-4R, IL-7R, and IL-9R, like IL-2R β , all associate with JAK-1. We have recently shown that IL-15 also activates JAK-1 and JAK-3. The activation of JAK-1 and JAK-3 presumably is vital to signal transduction mediated by IL-2, IL-4, IL-7, IL-9 and IL-15. However, it is clear that the distinct signals transduced by different γ_c users cannot be explained solely by JAK-1 and JAK-3. Unique actions, such as the induction of tyrosine phosphorylation of IRS-1 by IL-4, presumably play major

roles in determining cytokine-specific actions, and may reflect the abilities of specific receptor complexes to recruit different substrates for the activated JAKs or other kinases.

Finally, we contemplate that the discoveries described above can be used to design assays for identifying compounds that inhibit IL-2 dependent signaling. Specifically, we contemplate that compounds that disrupt the interaction between JAK-1 and the IL-2R β subunit, or between JAK-3 and the γ_c subunit of the IL-2 receptor will be potential inhibitors of IL-2 signalling. The descriptions provided in the foregoing Examples detail how interactions between the JAK kinases and the IL-2R β and γ_c chains of the receptor can be detected. Contemplated inhibitors will be identified by virtue of inhibiting these interactions.

To illustrate how an inhibitor can be identified according to the invented method, we have conducted experiments using a synthetic peptide as a model inhibitor to inhibit the interaction between γ_c and JAK-3 in vitro. Specifically, the peptide used in this procedure had an amino acid sequence corresponding to a portion of the γ_c chain that binds the JAK-3 protein. The assay employed standard immunoprecipitation and Western blotting methods. In the following exemplary case, the γ_c synthetic peptide prevented binding of wild type γ_c to JAK-3. Since the anti- γ_c antibody used in the immunoprecipitation procedure did not immunoprecipitate the complex between the γ_c synthetic peptide and the JAK-3 protein, inhibition of γ_c binding to JAK-3 was determined by virtue of the ability of the synthetic peptide to inhibit coprecipitation of the complex containing γ_c and JAK-3. Significantly, the fact that a negative control peptide failed to inhibit the interaction between γ_c and JAK-3 demonstrated the specificity of the inhibition. Thus, the assay described below clearly distinguished between agents that did and agents that did not inhibit the γ_c -JAK-3 interaction.

Example 15 describes an assay that can be used to identify compounds that inhibit IL-2 receptor activation by preventing the interaction between JAK-3 and the γ_c chain.

EXAMPLE 15

An Assay for Compounds that Inhibit IL-2 Receptor Complex Formation

YT cells or COS-7 cells expressing γ_c and JAK-3 were stimulated or not stimulated with IL-2 and then lysed with 10 mM Tris (pH 7.5) containing 2 mM EDTA, 0.15 M NaCl, 0.875% Brij 96, 0.125% NONIDET P40TM detergent, 0.4 mM sodium vanadate, 1 mM 4-(2-aminoethyl)-benzenesulfonyl fluoride hydrochloride (ICN), 2.5 mM leupeptin, 2.5 mM aprotinin. The peptide to be tested as an inhibitor corresponded to the first 19 amino acids of the γ_c cytoplasmic tail (sequence ERTMPRIPTLKNLEDLVTE) (SEQ ID NO:7). A negative control peptide that was not expected to inhibit the interaction between γ_c and JAK-3 corresponded to a tyrosine phosphorylated region of γ_c . The negative control peptide had the sequence APPCYTLKPET (SEQ ID NO:10), and was synthesized so that the Tyr residue at position 5 of the peptide was phosphotyrosine. The test peptide or the negative control peptide were separately added to the cell lysates to final concentrations of 140 μ M. The positive control immunoprecipitation was carried out in the absence of added peptide. Lysates were immunoprecipitated for 16 hours at 4° C. using antibodies to γ_c and protein A sepharose and washed six times before electrophoresis on SDS gels. The gels were immunoblotted with antisera to JAK-3 using ECL as described previously.

19

Results of the experiment described above indicated that only the competitor peptide partially inhibited γ_c -JAK-3 coprecipitation. As expected, and consistent with the findings presented under Example 12, only a weak association between the γ_c and JAK-3 proteins was observed in the absence of IL-2 stimulation.

In particular, the lane corresponding to the positive control coprecipitation in the extracts from IL-2 induced cells displayed an intense band on the x-ray film at a position corresponding to a molecular weight of 125 kDa. This result indicated the position of the JAK-3 protein on the immunoblot and provided a baseline for quantitative comparison with the test and negative control samples. The lane corresponding to the negative control sample had an intensity that was essentially identical to the band intensity in the positive control lane. This result indicated that the negative control peptide failed to inhibit association of γ_c and JAK-3 proteins. The lane corresponding to the sample containing the polypeptide being tested as an inhibitor had a band representing the JAK-3 protein that was of significantly diminished intensity. This result indicated that the 19 amino acid polypeptide having the sequence of SEQ ID NO:6 inhibited the γ_c -JAK-3 interaction. These findings illustrate the results that would be expected in assays for identifying inhibitors of the γ_c -JAK-3 interaction. In particular, an inhibitor of the γ_c -JAK-3 interaction can be identified by virtue of its ability to lessen the amount of the γ_c -JAK-3 coprecipitate compared to a trial that was conducted in the absence of any inhibitor.

20

Although the foregoing procedure was carried out by first immunoprecipitating with an anti- γ_c antibody and then Western blotting and probing with an anti-JAK-3 antibody, we anticipate the order of antibody usage could be reversed with equally good results. In particular, we anticipate that immunoprecipitation could be performed using the anti-JAK-3 antibody, and the Western blot probed with the anti- γ_c antibody. In this latter case, inhibitors of the γ_c -JAK-3 interaction would still be identified by virtue of diminishing the amount of γ_c -JAK-3 coprecipitate. The assay for inhibitors will involve identifying agents that cause a reduction in the amount of γ_c detected on the x-ray film relative to a trial that omitted the inhibitor.

Further, we anticipate that assays such as that described under Example 15 will be useful in the discovery of non-polypeptide drugs that inhibit the γ_c -JAK-3 interaction. These drugs will similarly be identified in such assays by virtue of their abilities to inhibit coprecipitation of the γ_c -JAK-3 complex.

Still further, we contemplate coprecipitation and Western blotting assays to identify compounds that inhibit the interaction between JAK-1 or JAK-3 and IL-2R β . Such assays can be readily performed using techniques well known to those having ordinary skill in the art. Inhibitors of this interaction are believed useful as drugs to inhibit transmembrane signaling that is dependent on the interaction between these receptor subunits. Specifically, these drugs are anticipated for use as immunomodulators which inhibit IL-2 dependent signaling.

SEQUENCE LISTING

<160> NUMBER OF SEQ ID NOS: 11

<210> SEQ ID NO 1

<211> LENGTH: 6

<212> TYPE: PRT

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: n-terminal fragment

<400> SEQUENCE: 1

Asp Val Trp Ser Phe Gly
1 5

<210> SEQ ID NO 2

<211> LENGTH: 28

<212> TYPE: DNA

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: synthetic primer

<220> FEATURE:

<221> NAME/KEY: misc_feature

<222> LOCATION: 14, 20, 26

<223> OTHER INFORMATION: n = A,T,C or G

<220> FEATURE:

<221> NAME/KEY: misc_feature

<222> LOCATION: 14, 20, 26

<223> OTHER INFORMATION: n = A,T,C or G

<400> SEQUENCE: 2

ccagcggccg cgtncaycgn gayctngc

28

<210> SEQ ID NO 3

<211> LENGTH: 28

<212> TYPE: DNA

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

-continued

<div><223> OTHER INFORMATION: synthetic primer</div> <div><220> FEATURE:</div> <div><221> NAME/KEY: misc_feature</div> <div><222> LOCATION: 17, 23</div> <div><223> OTHER INFORMATION: n = A,T,C or G</div> <div><220> FEATURE:</div> <div><221> NAME/KEY: misc_feature</div> <div><222> LOCATION: 17, 23</div> <div><223> OTHER INFORMATION: n = A,T,C or G</div> <div><400> SEQUENCE: 3</div> <div>ccagcggccg cccraanswc canacrtc</div>		28		
<div><210> SEQ ID NO 4</div> <div><211> LENGTH: 6</div> <div><212> TYPE: PRT</div> <div><213> ORGANISM: Artificial Sequence</div> <div><220> FEATURE:</div> <div><223> OTHER INFORMATION: n-terminal fragment</div> <div><220> FEATURE:</div> <div><221> NAME/KEY: VARIANT</div> <div><222> LOCATION: 2, 4, 5</div> <div><223> OTHER INFORMATION: Xaa = Any Amino Acid</div> <div><220> FEATURE:</div> <div><221> NAME/KEY: VARIANT</div> <div><222> LOCATION: 2, 4, 5</div> <div><223> OTHER INFORMATION: Xaa = Any Amino Acid</div> <div><400> SEQUENCE: 4</div> <div>Gly Xaa Gly Xaa Xaa Gly</div> <div>1 5</div>				
<div><210> SEQ ID NO 5</div> <div><211> LENGTH: 6</div> <div><212> TYPE: PRT</div> <div><213> ORGANISM: Artificial Sequence</div> <div><220> FEATURE:</div> <div><223> OTHER INFORMATION: n-terminal fragment</div> <div><400> SEQUENCE: 5</div> <div>Phe Trp Tyr Ala Pro Glu</div> <div>1 5</div>				
<div><210> SEQ ID NO 6</div> <div><211> LENGTH: 7</div> <div><212> TYPE: PRT</div> <div><213> ORGANISM: Artificial Sequence</div> <div><220> FEATURE:</div> <div><223> OTHER INFORMATION: n-terminal fragment</div> <div><400> SEQUENCE: 6</div> <div>Val Asp Gly Tyr Phe Arg Leu</div> <div>1 5</div>				
<div><210> SEQ ID NO 7</div> <div><211> LENGTH: 19</div> <div><212> TYPE: PRT</div> <div><213> ORGANISM: Artificial Sequence</div> <div><220> FEATURE:</div> <div><223> OTHER INFORMATION: n-terminal fragment</div> <div><400> SEQUENCE: 7</div> <div>Glu Arg Thr Met Pro Arg Ile Pro Thr Leu Lys Asn Leu Glu Asp Leu</div> <div>1 5 10 15</div>				
Val Thr Glu				
<div><210> SEQ ID NO 8</div> <div><211> LENGTH: 4064</div> <div><212> TYPE: DNA</div>				

-continued

<213> ORGANISM: Homo sapiens		
<220> FEATURE:		
<221> NAME/KEY: CDS		
<222> LOCATION: (96)...(3467)		
<400> SEQUENCE: 8		
ccctctgacc aggactgagg ggctttttct ctctgtgccc caggcaagtt gcactcatta	60	
tggaattccg gcggcccgct aggcaagttg cactc atg gca cct cca agt gaa	113	
Met Ala Pro Pro Ser Glu		
1 5		
gag acg ccc ctg atc cct cag cgt tca tgc agc ctc ttg tcc acg gag	161	
Glu Thr Pro Leu Ile Pro Gln Arg Ser Cys Ser Leu Leu Ser Thr Glu		
10 15 20		
gct ggt gcc ctg cat gtg ctg ctg ccc gct cgg gcc ccg ggg ccc ccc	209	
Ala Gly Ala Leu His Val Leu Leu Pro Ala Arg Ala Pro Gly Pro Pro		
25 30 35		
cag cgc cta tct ttc tcc ttt ggg gac cac ttg gct gag gac ctg tgc	257	
Gln Arg Leu Ser Phe Ser Phe Gly Asp His Leu Ala Glu Asp Leu Cys		
40 45 50		
gtg cag gct gcc aag gcc agc ggc atc ctg cct gtg tac cac tcc ctc	305	
Val Gln Ala Ala Lys Ala Ser Gly Ile Leu Pro Val Tyr His Ser Leu		
55 60 65 70		
ttt gct ctg gcc acg gag gac ctg tcc tgc tgg ttc ccc ccg agc cac	353	
Phe Ala Leu Ala Thr Glu Asp Leu Ser Cys Trp Phe Pro Pro Ser His		
75 80 85		
atc ttc tcc gtg gag gat gcc agc acc caa gtc ctg ctg tac agg att	401	
Ile Phe Ser Val Glu Asp Ala Ser Thr Gln Val Leu Leu Tyr Arg Ile		
90 95 100		
cgc ttt tac ttc ccc aat tgg ttt ggg ctg gag aag tgc cac cgc ttc	449	
Arg Phe Tyr Phe Pro Asn Trp Phe Gly Leu Glu Lys Cys His Arg Phe		
105 110 115		
ggg cta cgc aag gat ttg gcc agt gct atc ctt gac ctg cca gtc ctg	497	
Gly Leu Arg Lys Asp Leu Ala Ser Ala Ile Leu Asp Leu Pro Val Leu		
120 125 130		
gag cac ctc ttt gcc cag cac cgc agt gac ctg gtg agt ggg cgc ctc	545	
Glu His Leu Phe Ala Gln His Arg Ser Asp Leu Val Ser Gly Arg Leu		
135 140 145 150		
ccc gtg ggc ctc agt ctc aag gag cag ggt gag tgt ctc agc ctg gcc	593	
Pro Val Gly Leu Ser Leu Lys Glu Gln Gly Glu Cys Leu Ser Leu Ala		
155 160 165		
gtg ttg gac ctg gcc cgg atg gcg cga gag cag gcc cag cgg ccg gga	641	
Val Leu Asp Leu Ala Arg Met Ala Arg Glu Gln Ala Gln Arg Pro Gly		
170 175 180		
gag ctg ctg aag act gtc agc tac aag gcc tgc cta ccc cca agc ctg	689	
Glu Leu Leu Lys Thr Val Ser Tyr Lys Ala Cys Leu Pro Pro Ser Leu		
185 190 195		
cgc gac ctg atc cag ggc ctg agc ttc gtg acg cgg agg gct att cgg	737	
Arg Asp Leu Ile Gln Gly Leu Ser Phe Val Thr Arg Arg Ala Ile Arg		
200 205 210		
agg acg gtg cgc aga gcc ctg ccg cgc gtg gcc gcc tgc cag gca gac	785	
Arg Thr Val Arg Arg Ala Leu Pro Arg Val Ala Ala Cys Gln Ala Asp		
215 220 225 230		
cgg cac tcg ctc atg gcc aag tac atc atg gac ctg gag cgg ctg gat	833	
Arg His Ser Leu Met Ala Lys Tyr Ile Met Asp Leu Glu Arg Leu Asp		
235 240 245		
cca gcc ggg gcc gcc gag acc ttc cac gtg ggc ctc cct ggg gcc ctt	881	
Pro Ala Gly Ala Ala Glu Thr Phe His Val Gly Leu Pro Gly Ala Leu		
250 255 260		
ggt ggc cac gac ggg ctg ggg ctg ctc cgc gtg gct ggt gac ggc ggc	929	
Gly Gly His Asp Gly Leu Gly Leu Leu Arg Val Ala Gly Asp Gly Gly		
265 270 275		

-continued

atc gcc tgg acc cag gga gaa cag gag gtc ctc cag ccc ttc tgc gac Ile Ala Trp Thr Gln Gly Glu Gln Glu Val Leu Gln Pro Phe Cys Asp 280 285 290	977
ttt cca gaa atc gta gac att agc atc aag cag gcc ccg cgc gtt ggc Phe Pro Glu Ile Val Asp Ile Ser Ile Lys Gln Ala Pro Arg Val Gly 295 300 305 310	1025
ccg gcc gga gag cac cgc ctg gtc act gtt acc agg aca gac aac cag Pro Ala Gly Glu His Arg Leu Val Thr Val Thr Arg Thr Asp Asn Gln 315 320 325	1073
att tta gag gcc gag ttc cca ggg ctg ccc gag gct ctg tcg ttc gtg Ile Leu Glu Ala Glu Phe Pro Gly Leu Pro Glu Ala Leu Ser Phe Val 330 335 340	1121
gcg ctc gtg gac ggc tac ttc cgg ctg acc acg gac tcc cag cac ttc Ala Leu Val Asp Gly Tyr Phe Arg Leu Thr Thr Asp Ser Gln His Phe 345 350 355	1169
ttc tgc aag gag gtg gca ccg ccg agg ctg ctg gag gaa gtg gcc gag Phe Cys Lys Glu Val Ala Pro Pro Arg Leu Leu Glu Glu Val Ala Glu 360 365 370	1217
cag tgc cac ggc ccc atc act ctg gac ttt gcc atc aac aag ctc aag Gln Cys His Gly Pro Ile Thr Leu Asp Phe Ala Ile Asn Lys Leu Lys 375 380 385 390	1265
act ggg ggc tca cgt cct ggc tcc tat gtt ctc cgc cgc agc ccc cag Thr Gly Gly Ser Arg Pro Gly Ser Tyr Val Leu Arg Arg Ser Pro Gln 395 400 405	1313
gac ttt gac agc ttc ctc ctc act gtc tgt gtc cag aac ccc ctt ggt Asp Phe Asp Ser Phe Leu Leu Thr Val Cys Val Gln Asn Pro Leu Gly 410 415 420	1361
cct gat tat aag ggc tgc ctc atc cgg cgc agc ccc aca gga acc ttc Pro Asp Tyr Lys Gly Cys Leu Ile Arg Arg Ser Pro Thr Gly Thr Phe 425 430 435	1409
ctt ctg gtt ggc ctc agc cga ccc cac agc agt ctt cga gag ctc ctg Leu Leu Val Gly Leu Ser Arg Pro His Ser Ser Leu Arg Glu Leu Leu 440 445 450	1457
gca acc tgc tgg gat ggg ggg ctg cac gta gat ggg gtg gca gtg acc Ala Thr Cys Trp Asp Gly Gly Leu His Val Asp Gly Val Ala Val Thr 455 460 465 470	1505
ctc act tcc tgc tgt atc ccc aga ccc aaa gaa aag tcc aac ctg atc Leu Thr Ser Cys Cys Ile Pro Arg Pro Lys Glu Lys Ser Asn Leu Ile 475 480 485	1553
gtg gtc cag aga ggt cac agc cca ccc aca tca tcc ttg gtt cag ccc Val Val Gln Arg Gly His Ser Pro Pro Thr Ser Ser Leu Val Gln Pro 490 495 500	1601
caa tcc caa tac cag ctg agt cag atg aca ttt cac aag atc cct gct Gln Ser Gln Tyr Gln Leu Ser Gln Met Thr Phe His Lys Ile Pro Ala 505 510 515	1649
gac agc ctg gag tgg cat gag aac ctg ggc cat ggg tcc ttc acc aag Asp Ser Leu Glu Trp His Glu Asn Leu Gly His Gly Ser Phe Thr Lys 520 525 530	1697
att tac cgg ggc tgt cgc cat gag gtg gtg gat ggg gag gcc cga aag Ile Tyr Arg Gly Cys Arg His Glu Val Val Asp Gly Glu Ala Arg Lys 535 540 545 550	1745
aca gag gtg ctg ctg aag gtc atg gat gcc aag cac aag aac tgc atg Thr Glu Val Leu Leu Lys Val Met Asp Ala Lys His Lys Asn Cys Met 555 560 565	1793
gag tca ttc ctg gaa gca gcg agc ttg atg agc caa gtg tcg tac cgg Glu Ser Phe Leu Glu Ala Ala Ser Leu Met Ser Gln Val Ser Tyr Arg 570 575 580	1841
cat ctc gtg ctg ctc cac ggc gtg tgc atg gct gga gac agc acc atg His Leu Val Leu Leu His Gly Val Cys Met Ala Gly Asp Ser Thr Met 585 590 595	1889

-continued

gtg cag gaa ttt gta cac ctg ggg gcc ata gac atg tat ctg cga aaa Val Gln Glu Phe Val His Leu Gly Ala Ile Asp Met Tyr Leu Arg Lys 600 605 610	1937
cgt ggc cac ctg gtg cca gcc agc tgg aag ctg cag gtg gtc aaa cag Arg Gly His Leu Val Pro Ala Ser Trp Lys Leu Gln Val Val Lys Gln 615 620 625 630	1985
ctg gcc tac gcc ctc aac tat ctg gag gac aaa ggc ctg ccc cat ggc Leu Ala Tyr Ala Leu Asn Tyr Leu Glu Asp Lys Gly Leu Pro His Gly 635 640 645	2033
aat gtc tct gcc cgg aag gtg ctc ctg gct cgg gag ggg gct gat ggg Asn Val Ser Ala Arg Lys Val Leu Leu Ala Arg Glu Gly Ala Asp Gly 650 655 660	2081
agc ccg ccc ttc atc aag ctg agt gac cct ggg gtc agc ccc gct gtg Ser Pro Pro Phe Ile Lys Leu Ser Asp Pro Gly Val Ser Pro Ala Val 665 670 675	2129
tta agc ctg gag atg ctc acc gac agg atc ccc tgg gtg gcc ccc gag Leu Ser Leu Glu Met Leu Thr Asp Arg Ile Pro Trp Val Ala Pro Glu 680 685 690	2177
tgt ctc cgg gag gcg cag aca ctt agc ttg gaa gct gac aag tgg ggc Cys Leu Arg Glu Ala Gln Thr Leu Ser Leu Glu Ala Asp Lys Trp Gly 695 700 705 710	2225
ttc ggc gcc acg gtc tgg gaa gtg ttt agt ggc gtc acc atg ccc atc Phe Gly Ala Thr Val Trp Glu Val Phe Ser Gly Val Thr Met Pro Ile 715 720 725	2273
agt gcc ctg gat cct gct aag aaa ctc caa ttt tat gag gac cgg cag Ser Ala Leu Asp Pro Ala Lys Lys Leu Gln Phe Tyr Glu Asp Arg Gln 730 735 740	2321
cag ctg ccg gcc ccc aag tgg aca gag ctg gcc ctg ctg att caa cag Gln Leu Pro Ala Pro Lys Trp Thr Glu Leu Ala Leu Leu Ile Gln Gln 745 750 755	2369
tgc atg gcc tat gag ccg gtc cag agg ccc tcc ttc cga gcc gtc att Cys Met Ala Tyr Glu Pro Val Gln Arg Pro Ser Phe Arg Ala Val Ile 760 765 770	2417
cgt gac ctc aat agc ctc atc tct tca gac tat gag ctc ctc tca gac Arg Asp Leu Asn Ser Leu Ile Ser Ser Asp Tyr Glu Leu Leu Ser Asp 775 780 785 790	2465
ccc aca cct ggt gcc ctg gca cct cgt gat ggg ctg tgg aat ggt gcc Pro Thr Pro Gly Ala Leu Ala Pro Arg Asp Gly Leu Trp Asn Gly Ala 795 800 805	2513
cag ctc tat gcc tgc caa gac ccc acg atc ttc gag gag aga cac ctc Gln Leu Tyr Ala Cys Gln Asp Pro Thr Ile Phe Glu Glu Arg His Leu 810 815 820	2561
aag tac atc tca cag ctg ggc aag ggc aac ttt ggc agc gtg gag ctg Lys Tyr Ile Ser Gln Leu Gly Lys Gly Asn Phe Gly Ser Val Glu Leu 825 830 835	2609
tgc cgc tat gac ccg cta gcc cac aat aca ggt gcc ctg gtg gcc gtg Cys Arg Tyr Asp Pro Leu Ala His Asn Thr Gly Ala Leu Val Ala Val 840 845 850	2657
aaa cag ctg cag cac agc ggg cca gac cag cag agg gac ttt cag cgg Lys Gln Leu Gln His Ser Gly Pro Asp Gln Gln Arg Asp Phe Gln Arg 855 860 865 870	2705
gag att cag atc ctc aaa gca ctg cac agt gat ttc att gtc aag tat Glu Ile Gln Ile Leu Lys Ala Leu His Ser Asp Phe Ile Val Lys Tyr 875 880 885	2753
cgt ggt gtc agc tat ggc ccg ggc cgg cca gag ctg cgg ctg gtc atg Arg Gly Val Ser Tyr Gly Pro Gly Arg Pro Glu Leu Arg Leu Val Met 890 895 900	2801
gag tac ctg ccc agc ggc tgc ttg cgc gac ttc ctg cag cgg cac cgc Glu Tyr Leu Pro Ser Gly Cys Leu Arg Asp Phe Leu Gln Arg His Arg 905 910 915	2849

-continued

gcg cgc ctc gat gcc agc cgc ctc ctt ctc tat tcc tcg cag atc tgc Ala Arg Leu Asp Ala Ser Arg Leu Leu Leu Tyr Ser Ser Gln Ile Cys 920 925 930	2897
aag ggc atg gag tac ctg ggc tcc cgc cgc tgc gtg cac cgc gac ctg Lys Gly Met Glu Tyr Leu Gly Ser Arg Arg Cys Val His Arg Asp Leu 935 940 945 950	2945
gcc gcc cga aac atc ctc gtg gag agc gag gca cac gtc aag atc gct Ala Ala Arg Asn Ile Leu Val Glu Ser Glu Ala His Val Lys Ile Ala 955 960 965	2993
gac ttc ggc cta gct aag ctg ctg ccg ctt gac aaa gac tac tac gtg Asp Phe Gly Leu Ala Lys Leu Leu Pro Leu Asp Lys Asp Tyr Tyr Val 970 975 980	3041
gtc cgc gag cca ggc cag agc ccc att ttc tgg tat gcc ccc gaa tcc Val Arg Glu Pro Gly Gln Ser Pro Ile Phe Trp Tyr Ala Pro Glu Ser 985 990 995	3089
ctc tcg gac aac atc ttc tct cgc cag tca gac gtc tgg agc ttc ggg Leu Ser Asp Asn Ile Phe Ser Arg Gln Ser Asp Val Trp Ser Phe Gly 1000 1005 1010	3137
gtc gtc ctg tac gag ctc ttc acc tac tgc gac aaa agc tgc agc ccc Val Val Leu Tyr Glu Leu Phe Thr Tyr Cys Asp Lys Ser Cys Ser Pro 1015 1020 1025 1030	3185
tcg gcc gag ttc ctg cgg atg atg gga tgt gag cgg gat gtc ccc gcc Ser Ala Glu Phe Leu Arg Met Met Gly Cys Glu Arg Asp Val Pro Ala 1035 1040 1045	3233
ctc tgc cgc ctc ttg gaa ctg ctg gag gag ggc cag agg ctg ccg gcg Leu Cys Arg Leu Leu Glu Leu Leu Glu Gly Gln Arg Leu Pro Ala 1050 1055 1060	3281
cct cct gcc tgc cct gct gag gtt cac gag ctc atg aag ctg tgc tgg Pro Pro Ala Cys Pro Ala Glu Val His Glu Leu Met Lys Leu Cys Trp 1065 1070 1075	3329
gcc cct agc cca cag gac cgg cca tca ttc agc gcc ctg ggc ccc cag Ala Pro Ser Pro Gln Asp Arg Pro Ser Phe Ser Ala Leu Gly Pro Gln 1080 1085 1090	3377
ctg gac atg ctg tgg agc gga agc cgg ggg tgt gag act cat gcc ttc Leu Asp Met Leu Trp Ser Gly Ser Arg Gly Cys Glu Thr His Ala Phe 1095 1100 1105 1110	3425
act gct cac cca gag ggc aaa cac cac tcc ctg tcc ttt tca Thr Ala His Pro Glu Gly Lys His His Ser Leu Ser Phe Ser 1115 1120	3467
tagctcctgc ccgcagacct ctggattagg tctctgttga ctggctgtgt gaccttaggc	3527
ccggagctgc ccctctctgg gcctcagagg ccttatgagg gtcctctact tcaggaacac	3587
ccccatgaca ttgcatttgg gggggctccc gtggcctgta gaatagcctg tggcctttgc	3647
aatttgttaa ggttcaagac agatgggcat atgtgtcagt ggggctctct gagtcctggc	3707
ccaaagaagc aaggaaccaa atttaagact ctgcacatct cccaaccct taagccctgg	3767
ccccctgagt ttccttttct cgtctctctc tttttatttt ttttattttt atttttattt	3827
ttgagacaga gcctcgctcg ttaccacaggg tggagtgcag tggtagcgat ctcggtcac	3887
agtgcaacct ctgcttcca ggttcaageg attctcctgc ctacgcctcc cgagtagctg	3947
ggattacagg tgtgcaccac cacaccggc taattttttt tatttttaat agagatgagg	4007
tttcaccatg atggccaggc tgatctcgaa ctctaacct caagtgatcc tcccacc	4064
<210> SEQ ID NO 9	
<211> LENGTH: 1124	
<212> TYPE: PRT	
<213> ORGANISM: Homo sapiens	
<400> SEQUENCE: 9	

-continued

Met	Ala	Pro	Pro	Ser	Glu	Glu	Thr	Pro	Leu	Ile	Pro	Gln	Arg	Ser	Cys
1				5					10					15	
Ser	Leu	Leu	Ser	Thr	Glu	Ala	Gly	Ala	Leu	His	Val	Leu	Leu	Pro	Ala
			20					25					30		
Arg	Ala	Pro	Gly	Pro	Pro	Gln	Arg	Leu	Ser	Phe	Ser	Phe	Gly	Asp	His
		35					40					45			
Leu	Ala	Glu	Asp	Leu	Cys	Val	Gln	Ala	Ala	Lys	Ala	Ser	Gly	Ile	Leu
	50					55					60				
Pro	Val	Tyr	His	Ser	Leu	Phe	Ala	Leu	Ala	Thr	Glu	Asp	Leu	Ser	Cys
65					70					75					80
Trp	Phe	Pro	Pro	Ser	His	Ile	Phe	Ser	Val	Glu	Asp	Ala	Ser	Thr	Gln
				85					90					95	
Val	Leu	Leu	Tyr	Arg	Ile	Arg	Phe	Tyr	Phe	Pro	Asn	Trp	Phe	Gly	Leu
			100					105					110		
Glu	Lys	Cys	His	Arg	Phe	Gly	Leu	Arg	Lys	Asp	Leu	Ala	Ser	Ala	Ile
		115					120					125			
Leu	Asp	Leu	Pro	Val	Leu	Glu	His	Leu	Phe	Ala	Gln	His	Arg	Ser	Asp
	130					135					140				
Leu	Val	Ser	Gly	Arg	Leu	Pro	Val	Gly	Leu	Ser	Leu	Lys	Glu	Gln	Gly
145					150					155					160
Glu	Cys	Leu	Ser	Leu	Ala	Val	Leu	Asp	Leu	Ala	Arg	Met	Ala	Arg	Glu
				165					170					175	
Gln	Ala	Gln	Arg	Pro	Gly	Glu	Leu	Leu	Lys	Thr	Val	Ser	Tyr	Lys	Ala
			180					185					190		
Cys	Leu	Pro	Pro	Ser	Leu	Arg	Asp	Leu	Ile	Gln	Gly	Leu	Ser	Phe	Val
		195					200					205			
Thr	Arg	Arg	Ala	Ile	Arg	Arg	Thr	Val	Arg	Arg	Ala	Leu	Pro	Arg	Val
	210					215					220				
Ala	Ala	Cys	Gln	Ala	Asp	Arg	His	Ser	Leu	Met	Ala	Lys	Tyr	Ile	Met
225					230					235					240
Asp	Leu	Glu	Arg	Leu	Asp	Pro	Ala	Gly	Ala	Ala	Glu	Thr	Phe	His	Val
				245					250					255	
Gly	Leu	Pro	Gly	Ala	Leu	Gly	Gly	His	Asp	Gly	Leu	Gly	Leu	Leu	Arg
			260					265					270		
Val	Ala	Gly	Asp	Gly	Gly	Ile	Ala	Trp	Thr	Gln	Gly	Glu	Gln	Glu	Val
		275					280					285			
Leu	Gln	Pro	Phe	Cys	Asp	Phe	Pro	Glu	Ile	Val	Asp	Ile	Ser	Ile	Lys
	290					295					300				
Gln	Ala	Pro	Arg	Val	Gly	Pro	Ala	Gly	Glu	His	Arg	Leu	Val	Thr	Val
305					310					315					320
Thr	Arg	Thr	Asp	Asn	Gln	Ile	Leu	Glu	Ala	Glu	Phe	Pro	Gly	Leu	Pro
				325					330					335	
Glu	Ala	Leu	Ser	Phe	Val	Ala	Leu	Val	Asp	Gly	Tyr	Phe	Arg	Leu	Thr
			340					345					350		
Thr	Asp	Ser	Gln	His	Phe	Phe	Cys	Lys	Glu	Val	Ala	Pro	Pro	Arg	Leu
		355					360					365			
Leu	Glu	Glu	Val	Ala	Glu	Gln	Cys	His	Gly	Pro	Ile	Thr	Leu	Asp	Phe
	370					375					380				
Ala	Ile	Asn	Lys	Leu	Lys	Thr	Gly	Gly	Ser	Arg	Pro	Gly	Ser	Tyr	Val
385					390					395					400
Leu	Arg	Arg	Ser	Pro	Gln	Asp	Phe	Asp	Ser	Phe	Leu	Leu	Thr	Val	Cys
				405					410					415	
Val	Gln	Asn	Pro	Leu	Gly	Pro	Asp	Tyr	Lys	Gly	Cys	Leu	Ile	Arg	Arg

-continued

420																425																430																															
Ser	Pro	Thr	Gly	Thr	Phe	Leu	Leu	Val	Gly	Leu	Ser	Arg	Pro	His	Ser																																																
435																440																445																															
Ser	Leu	Arg	Glu	Leu	Leu	Ala	Thr	Cys	Trp	Asp	Gly	Gly	Leu	His	Val																																																
450																455																460																															
Asp	Gly	Val	Ala	Val	Thr	Leu	Thr	Ser	Cys	Cys	Ile	Pro	Arg	Pro	Lys																																																
465																470																475																480															
Glu	Lys	Ser	Asn	Leu	Ile	Val	Val	Gln	Arg	Gly	His	Ser	Pro	Pro	Thr																																																
485																490																495																															
Ser	Ser	Leu	Val	Gln	Pro	Gln	Ser	Gln	Tyr	Gln	Leu	Ser	Gln	Met	Thr																																																
500																505																510																															
Phe	His	Lys	Ile	Pro	Ala	Asp	Ser	Leu	Glu	Trp	His	Glu	Asn	Leu	Gly																																																
515																520																525																															
His	Gly	Ser	Phe	Thr	Lys	Ile	Tyr	Arg	Gly	Cys	Arg	His	Glu	Val	Val																																																
530																535																540																															
Asp	Gly	Glu	Ala	Arg	Lys	Thr	Glu	Val	Leu	Leu	Lys	Val	Met	Asp	Ala																																																
545																550																555																560															
Lys	His	Lys	Asn	Cys	Met	Glu	Ser	Phe	Leu	Glu	Ala	Ala	Ser	Leu	Met																																																
565																570																575																															
Ser	Gln	Val	Ser	Tyr	Arg	His	Leu	Val	Leu	Leu	His	Gly	Val	Cys	Met																																																
580																585																590																															
Ala	Gly	Asp	Ser	Thr	Met	Val	Gln	Glu	Phe	Val	His	Leu	Gly	Ala	Ile																																																
595																600																605																															
Asp	Met	Tyr	Leu	Arg	Lys	Arg	Gly	His	Leu	Val	Pro	Ala	Ser	Trp	Lys																																																
610																615																620																															
Leu	Gln	Val	Val	Lys	Gln	Leu	Ala	Tyr	Ala	Leu	Asn	Tyr	Leu	Glu	Asp																																																
625																630																635																640															
Lys	Gly	Leu	Pro	His	Gly	Asn	Val	Ser	Ala	Arg	Lys	Val	Leu	Leu	Ala																																																
645																650																655																															
Arg	Glu	Gly	Ala	Asp	Gly	Ser	Pro	Pro	Phe	Ile	Lys	Leu	Ser	Asp	Pro																																																
660																665																670																															
Gly	Val	Ser	Pro	Ala	Val	Leu	Ser	Leu	Glu	Met	Leu	Thr	Asp	Arg	Ile																																																
675																680																685																															
Pro	Trp	Val	Ala	Pro	Glu	Cys	Leu	Arg	Glu	Ala	Gln	Thr	Leu	Ser	Leu																																																
690																695																700																															
Glu	Ala	Asp	Lys	Trp	Gly	Phe	Gly	Ala	Thr	Val	Trp	Glu	Val	Phe	Ser																																																
705																710																715																720															
Gly	Val	Thr	Met	Pro	Ile	Ser	Ala	Leu	Asp	Pro	Ala	Lys	Lys	Leu	Gln																																																
725																730																735																															
Phe	Tyr	Glu	Asp	Arg	Gln	Gln	Leu	Pro	Ala	Pro	Lys	Trp	Thr	Glu	Leu																																																
740																745																750																															
Ala	Leu	Leu	Ile	Gln	Gln	Cys	Met	Ala	Tyr	Glu	Pro	Val	Gln	Arg	Pro																																																
755																760																765																															
Ser	Phe	Arg	Ala	Val	Ile	Arg	Asp	Leu	Asn	Ser	Leu	Ile	Ser	Ser	Asp																																																
770																775																780																															
Tyr	Glu	Leu	Leu	Ser	Asp	Pro	Thr	Pro	Gly	Ala	Leu	Ala	Pro	Arg	Asp																																																
785																790																795																800															
Gly	Leu	Trp	Asn	Gly	Ala	Gln	Leu	Tyr	Ala	Cys	Gln	Asp	Pro	Thr	Ile																																																
805																810																815																															
Phe	Glu	Glu	Arg	His	Leu	Lys	Tyr	Ile	Ser	Gln	Leu	Gly	Lys	Gly	Asn																																																
820																825																830																															
Phe	Gly	Ser	Val	Glu	Leu	Cys	Arg	Tyr	Asp	Pro	Leu	Ala	His	Asn	Thr																																																
835																840																845																															

-continued

Gly	Ala	Leu	Val	Ala	Val	Lys	Gln	Leu	Gln	His	Ser	Gly	Pro	Asp	Gln
850						855					860				
Gln	Arg	Asp	Phe	Gln	Arg	Glu	Ile	Gln	Ile	Leu	Lys	Ala	Leu	His	Ser
865					870					875					880
Asp	Phe	Ile	Val	Lys	Tyr	Arg	Gly	Val	Ser	Tyr	Gly	Pro	Gly	Arg	Pro
				885					890					895	
Glu	Leu	Arg	Leu	Val	Met	Glu	Tyr	Leu	Pro	Ser	Gly	Cys	Leu	Arg	Asp
			900					905					910		
Phe	Leu	Gln	Arg	His	Arg	Ala	Arg	Leu	Asp	Ala	Ser	Arg	Leu	Leu	Leu
		915					920					925			
Tyr	Ser	Ser	Gln	Ile	Cys	Lys	Gly	Met	Glu	Tyr	Leu	Gly	Ser	Arg	Arg
	930					935					940				
Cys	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Val	Glu	Ser	Glu
945					950					955					960
Ala	His	Val	Lys	Ile	Ala	Asp	Phe	Gly	Leu	Ala	Lys	Leu	Leu	Pro	Leu
			965						970					975	
Asp	Lys	Asp	Tyr	Tyr	Val	Val	Arg	Glu	Pro	Gly	Gln	Ser	Pro	Ile	Phe
			980					985					990		
Trp	Tyr	Ala	Pro	Glu	Ser	Leu	Ser	Asp	Asn	Ile	Phe	Ser	Arg	Gln	Ser
		995					1000						1005		
Asp	Val	Trp	Ser	Phe	Gly	Val	Val	Leu	Tyr	Glu	Leu	Phe	Thr	Tyr	Cys
	1010					1015					1020				
Asp	Lys	Ser	Cys	Ser	Pro	Ser	Ala	Glu	Phe	Leu	Arg	Met	Met	Gly	Cys
1025					1030					1035					1040
Glu	Arg	Asp	Val	Pro	Ala	Leu	Cys	Arg	Leu	Leu	Glu	Leu	Leu	Glu	Glu
			1045						1050					1055	
Gly	Gln	Arg	Leu	Pro	Ala	Pro	Pro	Ala	Cys	Pro	Ala	Glu	Val	His	Glu
			1060					1065					1070		
Leu	Met	Lys	Leu	Cys	Trp	Ala	Pro	Ser	Pro	Gln	Asp	Arg	Pro	Ser	Phe
		1075					1080					1085			
Ser	Ala	Leu	Gly	Pro	Gln	Leu	Asp	Met	Leu	Trp	Ser	Gly	Ser	Arg	Gly
	1090					1095					1100				
Cys	Glu	Thr	His	Ala	Phe	Thr	Ala	His	Pro	Glu	Gly	Lys	His	His	Ser
1105					1110					1115					1120
Leu	Ser	Phe	Ser												

<210> SEQ ID NO 10
<211> LENGTH: 11
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: n-terminal fragment
<220> FEATURE:
<221> NAME/KEY: PHOSPHORYLATION
<222> LOCATION: 5

<400> SEQUENCE: 10

Ala Pro Pro Cys Tyr Thr Leu Lys Pro Glu Thr
1 5 10

<210> SEQ ID NO 11
<211> LENGTH: 21
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: synthetic peptide

-continued

<400> SEQUENCE: 11															
Gly	Cys	Glu	Thr	His	Ala	Phe	Thr	Ala	His	Pro	Glu	Gly	Lys	His	His
1				5					10					15	
Ser	Leu	Ser	Phe	Ser											
				20											

What is claimed is:
1. An isolated polynucleotide encoding human JAK-3 protein, said JAK-3 protein being a protein tyrosine kinase that includes an amino acid sequence Gly-Cys-Glu-Thr-His-Ala-Phe-Thr-Ala-His-Pro-Glu-Gly-Lys-His-His-Ser-Leu-Ser-Phe-Ser (amino acids 1104-1124 of SEQ ID NO:9), said JAK-3 protein having a molecular weight of approximately 125 kDa measurable by SDS polyacrylamide gel electrophoresis in an 8% polyacrylamide gel and Western blotting

using an antibody specific for said amino acid sequence, that has tandem non-identical catalytic domains, lacks SH2 or SH3 domains, and is expressed in NK cells and stimulated or transformed T cells, but not substantially in resting T cells.
2. The isolated polynucleotide of claim 1, wherein said polynucleotide has the nucleotide sequence of SEQ ID NO:8.

* * * * *